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(54) Title: **PEPTIDE-LIPID CONJUGATES, LIPOSOMES AND LIPOSOMAL DRUG DELIVERY**

(57) Abstract: Peptide-lipid conjugates are incorporated into liposomes so as to selectively destabilize the liposomes in the vicinity of target peptidase-secreting cells, and hence, to deliver the liposomes to the vicinity of the target cells, or directly into the cells. The liposomes can thus be used to treat mammals for diseases, disorders or conditions, e.g., tumors, microbial infection and inflammations, characterized by the occurrence of peptidase-secreting cells.



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PEPTIDE-LIPID CONJUGATES, LIPOSOMES
AND LIPOSOMAL DRUG DELIVERY

Cross-References to Related Applications.

5 This application is a continuation-in-part ("CIP") of our copending U.S. application, Serial No. 09/168,010, filed October 7, 1998, which is a division of U.S. application Serial No. 08/950,618, filed October 15, 1997, which is a CIP that claims the benefit of provisional application U.S. Serial No. 60/027,544, filed October 15, 1996.

 This application is also a CIP of our copending U.S. application Serial No.
10 09/032,059, filed February 27, 1998, which is a CIP that claims the benefit of provisional application U.S. Serial No. 60/039,183, filed February 27, 1997.

Field of the Invention

 Peptide-lipid conjugates are incorporated into liposomes so as to localize
15 delivery of the liposomes' contents to the vicinity of target cells.

Background of the Invention

 Liposomes have been widely used as carriers to deliver a variety of therapeutic and diagnostic agents into cells. Encapsulation of active agents in liposomes protects
20 the agents from premature degradation, and ameliorates side effects resulting from administration of the agents to animals (for a review, see, e.g., A. Bangham, 1992; M. Ostro, 1987; and, M. Ostro and P. Cullis, 1989). However, the efficiency of liposomal drug delivery has heretofore been constrained by the lack of a means of inducing liposomes to preferentially release their contents in the vicinity of, or into, target cells.
25 This invention provides such a means, by incorporating peptide-lipid conjugates into liposomes and then contacting cells with these liposomes.

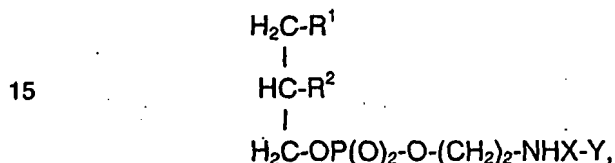
 The lipid portion of the peptide-lipid conjugate is a phosphatidylethanolamine ("PE"). Many of these lipids ordinarily do not organize into bilayers at neutral pH,
30 instead forming hexagonal (H_{II})-phase structures in aqueous environments which tend to destabilize the bilayers of liposomes into which the lipids have been incorporated.

- These same structures can also enhance the liposomes' fusogenicity (Verkleij, 1984; Cullis & de Kruijff, 1979; Ellens et al., 1989). Conjugation of a peptide to the PE stabilizes the PE in a bilayer conformation and hence, allows the conjugated lipid to be stably incorporated into liposome bilayers. However, once the peptide is cleaved, e.g.,
- 5 in the vicinity of peptidase-secreting cells, the lipid then resumes its nonbilayer-preferring conformation, in which it tends to destabilize the same liposome bilayers. In addition, the peptide linker could bind a blocking group, such as a polyethyleneglycol (PEG), such that it may act to sterically hinder fusion
- 10 The peptide portion of the peptide-lipid conjugate is any of those peptides having amino acid sequences that are recognized and cleaved by any of the various peptidases secreted by mammalian cells, e.g., at sites of inflammation and tumor metastases (see, e.g.: Aimes and Quigley, 1995; Fosang et al., 1994; Froelich et al., 1993; Knauper et al., 1996; Liotta et al., 1991; Moehrle et al., 1995; Nagase et al., 1994; Nakajima et al.,
- 15 1979; Otake et al., 1991; Palmieri et al., 1989; Pei et al., 1994; Prechel et al., 1995; Yamashita et al., 1994). Neither linkage of peptidase-cleavable peptides nor the incorporation of such peptides into liposomes, let alone for the purpose of promoting controlled liposome destabilization, has previously been described.
- 20 Vogel et al. (1993) and Subbaro et al. (1967) both covalently linked peptides to PEs; however, these peptide-lipids are not described therein as being cleavable by cell-secreted peptidases. Rather, the peptide-modified lipids of these documents are pH sensitive, adopting an alpha-helical conformation in low pH endosomal environments. Kirpotin et al. modified distearoyl phosphatidylcholine ("DSPE") by the attachment
- 25 thereto of methoxypoly(ethylene glycol) ("mPEG") to DSPE on the amino group; liposomes containing mPEG-modified DSPE were stable in solution until thiolytic cleavage and removal of the mPEG moiety. Kirpotin does not describe the peptide-based modification of PEs, let alone with peptidase-cleavable peptides.

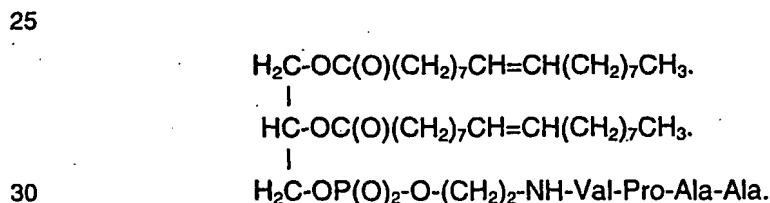
SUMMARY OF THE INVENTION

This invention provides a means of delivering and localizing the contents of liposomes to the vicinity of cells in a controlled manner, by conjugating certain peptides to phosphatidylethanolamines, and then incorporating these conjugated lipids into
 5 liposomes. The resulting liposomes are stable so long as the peptide remains conjugated to the lipid. However, once the peptide portion of the conjugate is cleaved from the lipid, by the action of cell-secreted or associated peptidases, the liposomes tend to destabilize, so as to release their contents in the vicinity of, or into, the secreting cells. Delivery of the liposomes' contents is thus targeted to the peptidase-secreting
 10 cells.

Peptide-lipid conjugates of this invention have the formula:



wherein: each of R^1 and R^2 is an acyl chain, X is a single bond or an acyl chain, and Y is
 20 a peptidase-cleavable peptide. The acyl chains are preferably oleic acid chains, X is preferably a single bond, and the peptide preferably contains the amino acid sequence Ala-Ala- (SEQ ID NO: 1); more preferably N-Acetyl-Ala-Ala- (SEQ ID NO: 2), or Ala-Ala-Pro-Val- (SEQ ID NO:3); and most preferably, N-methoxysuccinyl-Ala-Ala-Pro-Val (SEQ ID NO:4). Accordingly, the peptide-lipid conjugate preferably has the formula:



The liposomes' lipid component can be entirely composed of the peptide-lipid conjugate, or can comprise one or more additional lipids. Such additional lipids include,
 35 without limitation, any of the types of lipids, e.g., phospholipids, glycolipids and sterols,

which may be used in the preparation of liposomes. Most preferably, the liposome of this invention comprises a peptide-lipid conjugate and the positively charged synthetic lipid 1-N,N-dimethylamino dioleoyl propane (DODAP).

- 5 Controlled delivery with the liposomes of this invention can be used to deliver the liposomal drugs and/or bioactive agents in vitro or in vivo, for example, in the treatment of mammals afflicted with various diseases, disorders or conditions, e.g., cancers, amenable to treatment with the bioactive agent associated with the liposome.

10 **BRIEF DESCRIPTION OF THE DRAWINGS**

Figure 1. Structure of *N*-Ac-AA-DOPE and postulated scheme of conversion to DOPE by enzymatic cleavage.

- Figure 2.** TLC determination of protease mediated cleavage of *N*-Ac-AA-DOPE. *N*-Ac-AA-DOPE SUVs were incubated with elastase or proteinase K (1 mg enzyme/100 nmol lipid/0.1 ml buffer) overnight at 37°C (results of elastase-mediated cleavage are depicted in Fig. 2A, and results of proteinase K-mediated cleavage are depicted in Fig. 2B). Lipid was collected and separated by TLC. Lipid spots were developed as described hereinbelow. Lane 1, *N*-Ac-AA-DOPE without enzyme; lane 2, *N*-Ac-AA-DOPE with enzyme treatment; lane 3, DOPE from stock solution.

Figure 3. ³¹P NMR spectrum of *N*-Ac-alanyl-alanyl-dioleoyl phosphatidylethanolamine, DOPE and enzyme-treated liposomes.

Panel A., Pure *N*-Ac-ala-ala-DOPE

- 25 Panel B, Multilamellar liposomes, prepared as described below, comprising pure *N*-Ac-ala-ala-DOPE (2 umole total phospholipid) incubated in 200 ul of 154 mM NaCl, 10 mM TES, 0.1 mM EDTA at pH 7.4 and 37 degrees C for a total of two hours in the presence of 1 mg of proteinase K, as described in Example 2; and
Panel C, pure DOPE.

30

Figure 4. Proteinase K mediated cleavage of *N*-Ac-AA-DOPE. DOTAP/*N*-Ac-AA-DOPE (1:1) SUVs were incubated with or without elastase, proteinase K, or heat

inactivated proteinase K (95°C, 1 hour) at a concentration of 1 mg protease/100nmol lipid/0.1 ml buffer overnight at 37°C. Lipid was collected and analyzed by HPLC. The N-Ac-AA-DOPE peak was quantitated and the amount of cleavage was calculated as a percentage of the starting lipid. The bar indicates the range of data from several experiments.

Figure 5. TLC determination of the cleavage of MeO-suc-ala-ala-pro-val-DOPE. Fig. 5A. HLE (Human leukocyte elastase) dose titration : Lane 1/ 0 ug HLE/100nmol lipid; 2/ 5 ug HLE/100nmol lipid; 3/ 10 ug HLE/100nmol lipid; 4/ pure DOPE, 20ug; 5/ 20 ug HLE/100nmol lipid; 6/ 40 ug HLE/ 100nmol lipid; 7/ 40 ug proteinase K/100nmol lipid; and lane 8/ pure DOPE, 20ug;

Fig. 5B. Kinetics of HLE cleavage of MeO--suc-AAPV-PE 1/ without protease; 2/ 1 hour, 5 ug HLE/50nmol lipid; 3/ 2 hours, 5 ug HLE/50nmol lipid; 4/ pure DOPE, 20ug; 5/ 4 hours, 5 ug HLE/50nmol lipid; 6/ overnight, 5 ug HLE/50nmol lipid; 7/ pure DOPE, 20ug.

Fig. 5C Cleavage of MeO-suc-AAPV-PE by human neutrophil granule proteins: Lane 1/ without protease; 2/ 5 ug HLE/50nmol lipid; 3/ 2.5 ug granule proteins/100nmol lipid; 4/ pure DOPE, 20ug; 5/ 5 ug granule proteins/100nmol lipid; 6/ 10 ug granule proteins/100nmol lipid; 7/ 20 ug granule proteins/100nmol lipid; 8/ pure DOPE, 20ug.

Figure 6. Quantitation of elastase-mediated cleavage of MeO-suc-AAPV-DOPE to DOPE by ³¹P-NMR. DODAP/MeO-suc-AAPV-DOPE (1:1 mol/mol) freeze-thaw/extrusion vesicles were incubated with 0; 0.5; 1; 2; and 5 micrograms of elastase/100 nmol lipid. Samples were incubated for 2 hours at 37°C, after which liposomes were pelleted by ultracentrifugation. Liposomes were solubilized and monitored by ³¹P-NMR. Solid line: Percentage of total MEO-suc-AAPV-DOPE converted to DOPE; dotted line; % of expected MeO-suc-AAPV-DOPE on outer monolayers converted to DOPE assuming unilamellar vesicles, i.e., result multiplied by 2.

Figure 7. Effect of proteolytic enzymes on cleavage of N-Ac-ala-ala-PE. X-axis: % lipid cleaved; Y-axis: enzyme used (from top to bottom): S. caespitosus; S. griseus ("pronase"); peptidase; chymotrypsin; trypsin; protease type I; proteinase K; no enzyme added. Solid bars: experiment 1, using enzyme at a concentration of 0.5 mg

enzyme/100 nmole liposomes. Open bars: experiment 2, using enzyme at a concentration of 1 mg enzyme/10 nmole liposomes. Liposomes were sonicated SUVs composed of a 1:1 mixture of DOTAP/*N*-Ac-ala-ala-DOPE (molar ratio).

5 **Figure 8.** Determination of optimal liposomal composition. Liposomes were prepared having the molar ratios of DOTAP, *N*-Ac-AA-DOPE, PE, shown in the Figure. One mol % *N*-NBD-PE and *N*-Rho-PE fluorescent probes were included in all preparations. Liposomes were mixed with unlabeled PE/PS or PC/PS (80/20 mol%; 1:10 effector:acceptor ratio; 60uM total lipid) (Fig. 8A) or 2×10^8 RBC ghosts (Fig. 8B) at 37°C
10 for 1 hour. Lipid mixing was calculated as the percentage of *N*-NBD-PE FDQ relative to maximal FDQ, as determined by detergent addition. Binding of liposomes to RBC ghosts was quantitated after washing cells with buffer, by calculating the amount of *N*-Rho-PE fluorescence associated with the cell pellet relative to the total input fluorescence.

15 **Figure 9.** Concentration and time dependence of proteinase K activity. Activation of fusion: DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes (100nmol) containing fluorescent membrane probes were incubated in 0.1ml buffer at 37°C either overnight (Fig. 9A) with given amounts of proteinase K or (Fig. 9B) with 1 mg proteinase K for
20 given times. 10nmol aliquots were incubated with unlabeled PE/PS acceptor liposomes (80/20 mol%; 1:10 effector:acceptor ratio), after which lipid mixing was determined. *N*-Ac-AA-DOPE cleavage: unlabeled DOTAP/*N*-Ac-AA-DOPE (1:1 mol ratio) liposomes were treated identically as for fusion activation, after which lipid was extracted and analyzed by HPLC, and the amount of cleavage was calculated as described previously.

25

Figure 10. Elastase and proteinase K mediated activation of liposomal fusion. DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes containing fluorescent membrane probes were pretreated with human leukocyte elastase or proteinase K (1
30 mg protein/100 nmol lipid/0.1 ml buffer) overnight at 37°C. 10nmol aliquots were incubated with unlabeled PE/PS acceptor liposomes (80/20 mol%; 1:10

effector:acceptor ratio) for 60 min at 37°C. Lipid mixing was determined by monitoring *N*-NBD-PE FDQ.

- Figure 11.** Requirement for active proteinase K for DOTAP/*N*-Ac-AA-DOPE/PE liposome fusion activation with PS/PE liposomes. DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes (100nmol) containing fluorescent membrane probes were pretreated with or without 1 mg of proteinase K or heat inactivated proteinase K (1 hour, 95°C) overnight at 37°C in 0.1ml buffer. 10nmol aliquots were incubated with unlabeled PE/PS acceptor liposomes (80/20 mol%; 1:10 effector:acceptor ratio), after which lipid mixing was determined. Prot K carryover = effect of residual proteinase K carried over to incubation mixture with PE/PS liposomes was monitored by incubating untreated DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes with PE/PS liposomes in presence of freshly added proteinase K equivalent to the expected transferred amount.
- Figure 12.** Activation of DOTAP/*N*-Ac-AA-DOPE/PE liposomes by proteinase K for fusion with RBC ghosts. DOTAP/*N*-Ac-AA-DOPE/PE (20/10/70 mol%) liposomes (100nmol) containing fluorescent membrane probes were incubated overnight at 37°C with or without 1 mg of proteinase K in 0.1 ml buffer. 10nmol aliquots of DOTAP/*N*-Ac-AA-DOPE/PE liposomes as well as DOTAP/PE (20/80 mol%) liposomes were incubated with 1×10^8 RBC ghosts in buffer containing 0.5mM PMSF for 30min at 37°C, after which lipid mixing was determined. Effect of transferred proteinase K on lipid mixing was monitored by incubating untreated liposomes with RBC ghosts in presence of equivalent amount of proteinase K (prot K control).
- Figure 13** DOTAP/*N*-Ac-AA-DOPE/PE liposome with RBC ghosts: continuous kinetics of lipid mixing. 10nmol of DOTAP/*N*-Ac-AA-DOPE/PE (20/10/70 mol%) liposomes incubated (a) with or (b) without proteinase K overnight at 37°C were added to a cuvette containing 2 ml buffer with 0.5mM PMSF under continuous stirring and 37°C conditions. *N*-NBD-PE fluorescence recording was initiated and 1×10^8 RBC ghosts were added at 30 sec. (c) Effect of carryover proteinase K on lipid mixing was monitored by incubating untreated liposomes with RBC ghosts in presence of equivalent amount of proteinase K.

Figure 14 Dextran loaded DOTAP/N-Ac-AA-DOPE/PE liposome fusion with RBC ghosts. DOTAP/N-Ac-AA-DOPE/PE (20/10/70 mol%) liposomes were loaded with 10kD TX-red conjugated dextrans. Liposomes were incubated with proteinase K overnight at 37°C. 40nmol aliquots of dextran loaded liposomes or unloaded liposomes + free dextran were incubated with 1×10^8 RBC ghosts in 1 ml buffer for 30min at 37°C, after which cells were washed and observed by fluorescence microscopy or Nomarski differential interference contrast microscopy (Fig. 14A depicts results with dextran-loaded liposomes, using fluorescence microscopy; Fig. 14B depicts results with unloaded liposomes, using fluorescence microscopy; Fig. 14C. depicts results with dextran-loaded liposomes, using Nomarski contrast microscopy; Fig. 14D. depicts results with unloaded liposomes, using Nomarski contrast microscopy).

Figure 15 Optimum DODAP/MeO-suc-AAPV-DOPE liposome composition for elastase activation of binding/lipid mixing with HL60 cells. DODAP/MeO-suc-AAPV-DOPE liposomes prepared at 1:3, 1:1, or 3:1 (mol/mol) ratios and labeled with 0.75 mol% N-NBD-PE and 0.75 mol% N-Rho-PE were incubated with or without elastase ($5 \mu\text{g}$ elastase/100 nmol lipid) for 2 hours at 37°C. 10nmol of liposomes were then mixed with 1×10^6 HL60 cells and incubated for 30min at 37°C, pH 5. After washing twice with 5x volume of TES/NaCl/EDTA buffer, pH 7.4, A) binding and B) lipid mixing was determined by monitoring N-Rho-PE and N-NBD-PE fluorescence, respectively.

Figure 16. pH dependence of DODAP/MeO-suc-AAPV-DOPE liposome binding/lipid mixing with HL60 cells. Fluorescent lipid probe labeled DODAP/MeO-suc-AAPV-DOPE liposomes were incubated with or without elastase ($5 \mu\text{g}$ elastase/100 nmol lipid) for 2 hours at 37°C. 10nmol of liposomes were mixed with 1×10^6 HL60 cells in 200 μl TES/NaCl/EDTA buffer. Samples were incubated for 30min, 37°C, at the given pH and washed. Liposome binding (Fig. 16A) and lipid mixing (Fig. 16B) were determined by monitoring N-Rho-PE and N-NBD-PE fluorescence, respectively.

Figure 17 Active elastase is required for triggering DODAP/MeO-suc-AAPV-DOPE liposome binding/lipid mixing with HL60 cells. DODAP/MeO-suc-AAPV-DOPE liposomes containing 0.75 mol% N-NBD-PE and 0.75 mol% N-Rho-PE were incubated

alone, with elastase (5 • g elastase/100 nmol lipid), or with equivalent amount of heat inactivated elastase (95oC, 1 hour) for 2 hours at 37oC. 10nmol of pretreated liposomes were mixed with 1x10⁶ HL60 cells in 200 • l TES/NaCl/EDTA buffer. Samples were incubated for 30min, 37oC, at either pH 7.4 or pH 5 and washed.

- 5 Liposome binding (Fig. 17A) and lipid mixing (Fig. 17B) was determined as previously described.

Figure 18 Confocal microscopy of DODAP/MeO-suc-AAPV-DOPE liposome interaction with HL60 cells. DODAP/MeO-suc-AAPV-DOPE liposomes labeled with 0.75 mol% N-
10 NBD-PE and 0.75 mol% N-Rho-PE were incubated alone or with elastase (5 • g elastase/100 nmol lipid) for 2 hours at 37oC as described in Fig. 17. After washing, cells were observed for N-Rho-PE fluorescence. Color range employed as in Figure 19.

Figure 19 Delivery of TMR-dextran from DODAP/MeO-suc-AAPV-DOPE liposomes to
15 HL60 cells. DODAP/MeO-suc-AAPV-DOPE liposomes loaded with 10,000 MW TMR-dextran were incubated alone or with elastase (5 • g elastase/100 nmol lipid) for 2 hours at 37oC. Liposomes were then added directly to cells or subjected to 5 freeze/thaw cycles to release encapsulated TMR-dextran. 40 nmol of intact or freeze/thawed liposomes were mixed with 2x10⁵ HL60 cells and incubated for 30 min at 37oC, pH 5.
20 After washing, cells were observed for TMR-dextran fluorescence by confocal microscopy. Color range of 0-255 pixel values (0, bottom; 255, top) are given in upper right corner of each image.

Figure 20 Binding/lipid mixing of DODAP/MeO-suc-AAPV-DOPE liposomes with
25 ECV304 cells at pH 7.4. DODAP/MeO-suc-AAPV-DOPE liposomes containing N-NBD-PE and N-Rho-PE were incubated alone or with elastase (5 • g elastase/100 nmol lipid) for 2 hours at 37oC. 100nmol of pretreated liposomes were added to ECV304 cells that had been pretreated with biotinylated wheat germ agglutinin (WGA) and streptavidin as described in the Examples). After a 30min incubation at room temperature the cell
30 monolayer was washed to remove unbound liposomes and samples were incubated for additional 2 hours at 37oC. Binding and lipid mixing was determined by monitoring N-Rho-PE and N-NBD-PE fluorescence, respectively.

Figure 21 Confocal microscopy of elastase-activated calcein delivery from DODAP/MeO-suc-AAPV-DOPE liposomes to ECV304 cells at pH 7.4. DODAP/MeO-suc-AAPV-DOPE liposomes loaded with quenched concentration of calcein were

5 incubated alone or with elastase (5 • g elastase/100 nmol lipid) for 2 hours at 37oC. 100nmol of pretreated intact liposomes were added to ECV304 cells that had been pretreated with biotinylated WGA and streptavidin. After a 30 min incubation at room temperature the cell monolayer was washed to remove unbound liposomes and samples were incubated for an additional 4 hours at 37oC. Cells were observed by

10 confocal microscopy for calcein fluorescence. Color range of 0-255 pixel values (0, bottom; 255, top) are given in upper right corner of each image.

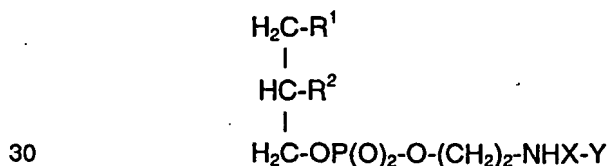
Figure 22 Elastase-triggered calcein delivery from DODAP/MeO-suc-AAPV-DOPE liposomes to ECV304 cells at pH 7.4. DODAP/MeO-suc-AAPV-DOPE liposomes loaded

15 with self-quenched concentration of calcein and N-Rho-PE as a lipid marker were incubated with or without elastase (5 • g elastase/100 nmol lipid) for 2 hours at 37oC. 100nmol liposomes were added to ECV304 cells that had been pretreated with biotinylated WGA and streptavidin. After a 30min incubation at room temperature the cell monolayer was washed to remove unbound liposomes and samples were incubated

20 for 0, 1, 2, 4, or 6 hours at 37oC. At given times calcein delivery was determined by monitoring calcein fluorescence.

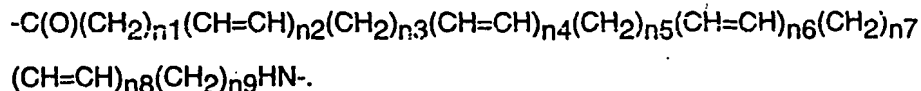
DETAILED DESCRIPTION OF THE INVENTION

25 This invention provides a peptide-lipid conjugate having the following formula:



wherein: each of R^1 and R^2 is independently a group having the formula
 $-\text{OC}(\text{O})(\text{CH}_2)_{n1}(\text{CH}=\text{CH})_{n2}(\text{CH}_2)_{n3}(\text{CH}=\text{CH})_{n4}(\text{CH}_2)_{n5}(\text{CH}=\text{CH})_{n6}(\text{CH}_2)_{n7}$

$(\text{CH}=\text{CH})_{n8}(\text{CH}_2)_{n9}\text{CH}_3$ and X is a linker moiety selected from the group consisting of a single bond and an acyl chain having the formula:



5

$n1$ is equal to zero or an integer of from 1 to 22, $n3$ is equal to zero or an integer of from 1 to 19, $n5$ is equal to zero or an integer of from 1 to 16, $n7$ is equal to zero or an integer of from 1 to 13 and $n9$ is equal to zero or an integer of from 1 to 10; and, each of $n2$, $n4$, $n6$ and $n8$ is independently zero or 1. For R^1 and R^2 , the sum of $n1 + 2n2 + n3 + 2n4 + n5 + 2n6 + n7 + 2n8 + n9$ is equal to an integer of from 10 to 22.

10

X is preferably a single bond; however, when X is other than a single bond, the sum of $n1 + 2n2 + n3 + 2n4 + n5 + 2n6 + n7 + 2n8 + n9$ for X is equal to an integer of from 1 to 22. X is then preferably saturated, most preferably:

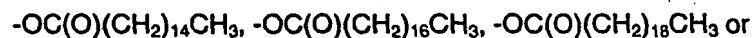


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Preferably, at least one of R^1 and R^2 contains at least one double bond, and the peptide-lipid conjugate is thus partially or completely unsaturated. More preferably, both of R^1 and R^2 contain one double bond, and the conjugate is thus completely unsaturated. Most preferably, presently, both R^1 and R^2 are

20

$-\text{OC}(\text{O})(\text{CH}_2)_7(\text{CH}=\text{CH})(\text{CH}_2)_7\text{CH}_3$, i.e., the peptide-lipid conjugate is a dioleoyl phosphatidylethanolamine (DOPE)-based conjugate. However, each of R^1 and R^2 can also be saturated or unsaturated acyl chains that include, without limitation:



25



Y is an "enzyme-cleavable peptide," which is a peptide comprising an amino acid sequence that is recognized by a peptidase expressed by a mammalian cell and found in surrounding tissue, or produced by a microbe capable of establishing an infection in a mammal. Enzyme-cleavable peptides can, but are not required to, contain one or more amino acids in addition to the amino acid recognition sequence; additional amino acids

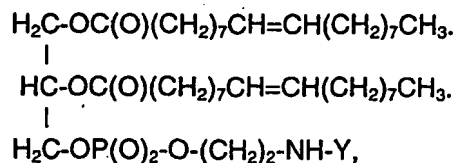
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or other groups (such as methoxysuccinyl, PEG, acetyl) can be added to the amino terminal, carboxy terminal, or both the amino and carboxy terminal ends of the recognition sequence. Means of adding amino acids to an amino acid sequence, e.g., in an automated peptide synthesizer, as well as means of detecting cleavage of a peptide by a peptidase, e.g., by chromatographic analysis for the amino acid products of such cleavage, are well known to ordinarily skilled artisans given the teachings of this invention.

Enzyme-cleavable peptides, typically from about 2 to 20 amino acids in length, are of sufficient length to project above the surfaces of lipid-based carriers into which they have been incorporated. Such peptides are well known to ordinarily skilled artisans given the teachings of this invention and include, for example and without limitation, the amino acid sequences: Ala-Ala-, Ala-Ala-Pro-Val, Ala-Ala-Met-, Ala-Ala-Pro-Phe-, Ala-Ala-Pro-Met-, Ala-Ala-Arg, Ser-Ala-Ala-Arg-, Ser-Ser-Ala-Ala-Arg-, Ser-S carboxyl sugar-Ala-Ala-Arg-, Ala-Ala-Asp-, Ser-Ala-Ala-Asp-, Ser-Ser-Ala-Ala-Asp-, Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ser-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva, Pro-Cha-Gly-Nva-His-Ala-Dpa-NH₂, Pro-Leu-Gly-Leu-Dpa-Ala-Arg-NH₂, Pro-Cha-Gly-Nva-, Pro-Leu-Gly-Leu, Gly-Pro-Arg, Leu-Pro-Arg, Glu-Gly-Arg, Pro-Leu-Gly-Leu- and Gly-Pro-Gln-Gly-Ile-. Presently, the preferred peptides comprise the amino acid sequence Ala-Ala, more preferably, Ala-Ala-Pro-Val; and most preferably, N-methoxysuccinyl-Ala-Ala-Pro-Val.

Accordingly, the peptide-lipid conjugate of this invention most preferably has the formula:

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wherein the peptide comprises the amino acid sequence N-methoxysuccinyl-Ala-Ala-Pro-Val.

Enzyme-cleavable peptides can be modified at their amino termini, for example, so as to increase their hydrophilicity, the stability of the liposome, or to enhance enzyme activity. Increased hydrophilicity enhances exposure of the peptides on the surfaces of lipid-based carriers into which the parent peptide-lipid conjugates have been

5 incorporated and can increase the liposome's stability. Groups suitable for attachment to peptides so as to increase their hydrophilicity are well known, and include, for example and without limitation: acetyl ("Ac"), acetyl-serine ("Ac-Ser"), acetyl-seryl-serine ("Ac-Ser-Ser-"), succinyl ("Suc"), succinyl-serine ("Suc-Ser"), succinyl-seryl-serine ("Suc-Ser-Ser-"), methoxy succinyl ("MeO-Suc"), methoxy succinyl-serine ("MeO-Suc-Ser-"),

10 methoxy succinyl-seryl-serine ("MeO-Suc-Ser-Ser-") and seryl-serine ("Ser-Ser-") groups, polyethylene glycol ("PEG"), polyacrylamide, polyacrylomorpholine, polyvinylpyrrolidine, a polyhydroxyl group and carboxy sugars, e.g., lactobionic, N-acetyl neuraminic and sialic acids, groups. The carboxy groups of these sugars would be linked to the N-terminus of the peptide via an amide linkage. Specific groups, such as acetyl, succinyl,

15 methoxysuccinyl or 3-cyclohexylalanyl ("Cha") groups, although not hydrophilic, enhance enzyme activity. Specific groups, such as polyethylene glycol, polyacrylamide, polyacrylomorpholine, polyvinylpyrrolidine, a polyhydroxyl group and carboxy sugars, e.g., lactobionic, N-acetylneuraminic and sialic acids, may additionally block exposure of the lipid bilayer to serum or other cellular components, prior to cleavage at the peptide-

20 linkage by a peptidase. Presently, the preferred N-terminal modification is an acetyl modification, and the more preferred modification is a methoxy-succinyl modification.

For purposes of the present invention, the term extracellular enzyme is defined to include enzymes that are produced by cells and that are released into the medium or

25 other areas surrounding the cell and is free from the cell; it is also meant to include enzymes which are produced by cells and secreted outside the cell, and which may remain in the immediate area of the cell, and also includes enzymes which are secreted by cells, which remain either bound to or associated with, the cell's outer surface. This definition is also meant to encompass the cell-secreted peptidases described below, and

30 enzymes produced by mammalian cells or by microbes.

Cell-secreted peptidases which recognize particular amino acid sequences are also well known to ordinarily skilled artisans given the teachings of this invention. Such peptidases include, for example and without limitation: matrix metalloproteinases, serine proteases, cysteine proteases, elastase, plasmin, plasminogen activators, such as tissue plasminogen activator and urokinase, stromelysin, human collagenases, cathepsins, lysozyme, granzymes, dipeptidyl peptidases, peptide hormone-inactivating enzymes, kininases, bacterial peptidases and viral proteases. Elastase, for example, is involved in tumor cell tissue remodeling; the breast cancer cell line MCF-7 has been shown to secrete elastase, the levels of which are inversely correlated to overall survival in breast cancer patients (Yamashita et al.). Moreover, the matrix metalloproteinase, stromelysin-3 ("ST3"), has been localized to the stromal area of tumor cells (Pei et al, 1994.); it specifically cleaves α_1 proteinase inhibitor between amino acids 350 and 351 (Ala-Met). Stromelysin-1 ("MMP-3") is also localized to areas of tissue remodeling, including sites of inflammation and tumor stroma (Nagase et al., 1994). Metastatic cancer cells display enhanced extracellular activity of, among others, the matrix metalloproteinases and urokinase-type plasminogen activator (reviewed in Liotta et al., 1991).

The cDNA of human collagenase-3 or MMP-13, another metalloproteinase was isolated from a breast tumor library (Knäuper et al., 1996); this enzyme cleaves peptides containing the amino acid sequences Pro-Cha-Gly-Nva-His- and Pro-Leu-Gly-Leu-. Furthermore, the 72 kDa gelatinase (MMP-2) is involved in regulating tumor cell invasiveness, and cleaves the amino acid sequence Gly-Pro-Gln-Gly-Ile- between the Gly and Ile residues (Aimes and Quigley, 1995; Liotta et al., 1991). Human neutrophils also secrete collagenases at sites of inflammation such as MMP-8 (neutrophil collagenase) and MMP-9 (type IV collagenase, 92 kDa gelatinase) (Fosang et al., 1994). Cathepsin G is also secreted from human neutrophils at sites of inflammation; its specificity is greatest for peptides containing the amino acid sequences Suc-Ala-Ala-Pro-Phe- or MeOSuc-Ala-Ala-Pro-Met- (Nakajima et al., 1979). Other enzymes secreted by neutrophils at sites of inflammation include cathepsins B and D as well as lysozyme. Granzymes A and B are secreted by cytotoxic lymphocytes in the synovial fluid of rheumatoid arthritis patients (Froehlich et al., 1993); granzyme A cleaves

peptides comprising Gly-Arg- and Ala-Ala-Arg- most efficiently, while granzyme B cleaves peptides comprising the amino acid sequence Ala-Ala-Asp (Otake et al., 1991).

Peptidases which hydrolyze enzyme-cleavable peptides also include the group of
5 enzymes that inactivate peptide hormones, e.g., aminopeptidase P and angiotensin-
converting enzyme, localized on the surface of endothelial cells. Aminopeptidase P
cleaves the Arg-Pro bond in bradykinin, and is localized to lung endothelial cells
(Prechel et al., 1995).

10 Numerous pathological conditions, some of which have already been discussed,
are associated with elevated enzyme activity. Metastatic cancer cells display enhanced
extracellular activity of several degradative enzymes, [for review see Liotta et al., 1991].
Elevated enzymatic activity facilitates the extravasation of these cells from the
circulation and increases their invasive potential. Other disorders, such as the
15 inflammatory conditions cystic fibrosis [McElvaney et al, 1991; Rees and Brain, 1995;
Berger et al, 1989, and Suter et al., 1986], rheumatoid arthritis [Al-haik et al, 1984, and
Gysen et al, 1985], and emphysema [Damiano et al, 1986, and Snider et al., 1991] are
accompanied by an increase in extracellular elastase activity due to release of elastase
from phagocytic cells. Elevated elastase activity appears to be due, in part, to an
20 imbalance in the elastase/anti-protease ratio [McElvaney et al., 1991, Cavarra et al.,
1996, and Doring et al., 1994]. Elastase has also been associated with tumor
progression and development [Yamashita et al., 1994, Yamashita et al., 1997, and
Starcher et al., 1996]. The ubiquitous yet specific nature of disease-associated
enzymatic activity, its localization near or on the membranes of cells involved in tissue
25 remodeling [Sato et al., 1994 and Owen et al., 1995] and its association with several
pathologies provide numerous opportunities for triggering specific liposomal delivery to
desired targets using the activity of such enzymes. The triggering event would be
expected to convert the liposome from a relatively inert state to a fusogenic state and
may even trigger specific binding depending on the design.

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The selectivity of liposomal activation can be modulated by the choice of an
enzyme substrate conjugated to a fusogenic lipid such that enzymatic cleavage releases

or unmasks fusogenic lipids. Thus liposomes may be designed for a selected site of activation and hence liposomal delivery could be targeted.

Peptide-lipid conjugates are prepared by any of a number of means for forming
5 an amide bond between the amino group of a phosphatidylethanolamine and the
carboxy terminus of an amino acid sequence. Such means include, without limitation,
those described in Example 1, hereinbelow. Briefly, an enzyme-cleavable peptide
containing an N-terminal blocking group is prepared as an anhydride; a
phosphatidylethanolamine such as DOPE is then reacted with the anhydride in the
10 presence of suitable reagents, such as triethylamine.

This invention also provides a liposome having a lipid component which
comprises the peptide-lipid conjugate of the invention. "Liposomes" are self-assembling
structures comprising one or more lipid bilayers, each of which surrounds an aqueous
15 compartment and comprises two opposing monolayers of amphipathic lipid molecules.
Amphipathic lipids comprise a polar (hydrophilic) headgroup region covalently linked to
one or two non-polar (hydrophobic) acyl chains. Energetically unfavorable contacts
between the hydrophobic acyl chains and the aqueous medium are generally believed to
induce lipid molecules to rearrange such that the polar headgroups are oriented towards
20 the aqueous medium while the acyl chains reorient towards the interior of the bilayer.
An energetically stable structure is formed in which the acyl chains are effectively
shielded from coming into contact with the aqueous medium.

Liposomes of this invention can have a single lipid bilayer (unilamellar
25 liposomes, "ULVs"), or multiple lipid bilayers (multilamellar liposomes, "MLVs"), and can
be made by a variety of methods well known in the art. These methods include without
limitation: Bangham's methods for making multilamellar liposomes (MLVs); Lenk's,
Fountain's and Cullis' methods for making MLVs with substantially equal interlamellar
solute distribution (see, for example, U.S. Patent Nos. 4,522,803, 4,588,578, 5,030,453,
30 5,169,637 and 4,975,282); and Papahadjopoulos et al.'s reverse-phase evaporation
method (U.S. Patent No. 4,235,871) for preparing oligolamellar liposomes. ULVs can

be produced from MLVs by such methods as sonication or extrusion (U.S. Patent No. 5,008,050 and U.S. Patent No. 5,059,421). The liposome of this invention can be produced by the methods of any of these disclosures, the contents of which are incorporated herein by reference.

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Various methodologies, such as sonication, homogenization, French Press application and milling can be used to prepare liposomes of a smaller size from larger liposomes. Extrusion (see U.S. Patent No. 5,008,050) can be used to size reduce liposomes, that is to produce liposomes having a predetermined mean size by forcing
10 the liposomes, under pressure, through filter pores of a defined, selected size. Tangential flow filtration (see WO89/008846), can also be used to regularize the size of liposomes, that is, to produce a population of liposomes having less size heterogeneity, and a more homogeneous, defined size distribution. The contents of these documents are incorporated herein by reference. Liposome sizes can also be determined by a
15 number of techniques, such as quasi-elastic light scattering, and with equipment, e.g., Nicomp® particle sizers, well within the possession of ordinarily skilled artisans.

Liposomes of this invention can have lipid components entirely composed of a peptide-lipid conjugate. However, the liposomes preferably contain one or more
20 additional lipids, including any of those lipids, such as phospholipids, glycolipids and sterols, typically used to prepare liposomes. Preferably, the additional lipid is a positively charged lipid, more preferably such a lipid selected from the group consisting of 1,2-bis(oleoyloxy)-3-(trimethyl ammonio)propane (DOTAP), 1-N,N-dimethylamino dioleoyl propane (DODAP), 1-oleoyl- 2-hydroxy-3-N,N-dimethylamino propane, 1,2-
25 diacyl-3-N,N-dimethylamino propane and 1,2-didecanoyl -1-N,N,-dimethylamino propane, 3 β -[N-[(N',N'-dimethylamino)ethane]carbamoyl]cholesterol (DC-Chol), 1,2-dimyristooxypropyl-3-dimethylhydroxyethyl ammonium bromide (DMRIE) and 1,2-dioleooxypropyl-3-dimethylhydroxyethyl ammonium bromide (DORI).

30 Most preferably, presently, the positively charged lipid is DODAP. Positively charged lipids are incorporated into the liposomes, preferably in at most about equimolar concentration respective to the peptide-lipid conjugate, in order to adjust the

net charge of the carrier. Increasing the positive charge on a lipid-based carrier enhances electrostatic interactions between the carrier and a biological membrane and hence, fusion between the carrier and the membrane.

5 The additional lipid can also include one or more phospholipids, such as a phosphatidylcholine ("PC"), which are generally added to lipid carriers to serve as structural stabilizers, or a phosphatidylethanolamine ("PE"). The PE may be selected from the group consisting of trans-esterified phosphatidylethanolamine (tPE), dipalmitoyl phosphatidylethanolamine (DPPE), palmitoyloleoyl phosphatidylethanolamine (POPE)
10 and dioleoyl phosphatidylethanolamine (DOPE); such additional PE's can be fusogenic because of the relatively unhydrated state of their headgroups.

 Alternatively, the PE is a PE to the headgroup of which is attached a moiety selected from the group consisting of dicarboxylic acids, polyethylene glycols, polyalkyl
15 ethers and gangliosides. Such modified PEs, also known as "headgroup-modified lipids," can inhibit the binding of serum proteins to lipid carriers such that the pharmacokinetic behavior of the carriers in the circulatory systems of animals is altered (see, e.g., Blume et al., 1993, Gabizon et al., 1993; Park et al., 1992, Woodle et al., and Allen et al.; the contents of which are incorporated herein by reference). The amount of
20 the headgroup-modified lipid incorporated into the liposomes depends upon a number of factors well known to the ordinarily skilled artisan, or within his purview to determine without undue experimentation. These include, but are not limited to: the type of lipid and the type of headgroup modification; the type and size of the liposome; and the intended therapeutic use of the formulation. The concentration of the headgroup-
25 modified lipid in the liposome is generally sufficient to prolong the liposome's circulatory half-life in an animal, but is not so great as induce unwanted side effects in the animal, and is typically at least about five mole percent of the lipid present in the liposome. Preferred headgroup-derivatized lipids include phosphatidylethanolamine-dicarboxylic acids ("PE-DCAs") and polyethyleneglycol- modified (PEGylated) lipids (for a
30 description of which, see Woodle et al. and Allen et al.).

The liposome of this invention can comprise a "targeting moiety," i.e., a moiety that can be attached to a liposome and which can then direct the liposome to a specific site within the body of a mammal. Such directed delivery is generally believed to occur as a result of the recognition by the targeting moiety of a compound on the surface of the cells being targeted. Typical targeting moieties include, without limitation, antibodies, cell receptor ligands, lectins and the like. Targeting moieties can be attached to liposomes by any of the means generally accepted in the art for the covalent or noncovalent attachment of such moieties to liposomes. Such means include, for example and without limitation, those described in the following documents, the contents of which are incorporated herein by reference: U.S. Patent No. 5,399,331 describes the coupling of proteins to liposomes through use of a crosslinking agent having at least one maleimido group and an amine reactive function; U.S. Patent Nos. 4,885,172, 5,059,421 and 5,171,578 link proteins to liposomes through use of the glycoprotein streptavidin; Sato and Sunamoto (1993) describe the coating of targeted liposomes with polysaccharides.

The liposomes of this invention can comprise one or more "bioactive agents," which are compounds or compositions of matter having biological, including therapeutic or diagnostic, activity in animals. Bioactive agents which may be associated with the liposomes include, but are not limited to: antiviral agents such as acyclovir, zidovudine and the interferons; antibacterial agents such as aminoglycosides, cephalosporins and tetracyclines; antifungal agents such as polyene antibiotics, imidazoles and triazoles; antimetabolic agents such as folic acid analogues, folic acid antagonists including methotrexate, and purine and pyrimidine analogs; antineoplastic agents such as the anthracycline antibiotics and plant alkaloids; sterols such as cholesterol; carbohydrates, e.g., sugars and starches; amino acids, peptides, proteins such as cell receptor proteins, immunoglobulins, enzymes, hormones, neurotransmitters and glycoproteins; dyes; radiolabels such as radioisotopes and radioisotope-labeled compounds; radiopaque compounds; fluorescent compounds; mydriatic compounds; bronchodilators; local anesthetics; nucleic acid sequences such as messenger RNA, cDNA, genomic DNA and plasmids; bioactive lipids such as ether lipids and ceramides; and the like.

Preferred bioactive agents are selected from the group consisting of nucleic acid sequences, antimicrobial agents, anticancer agents and anti-inflammatory agents.

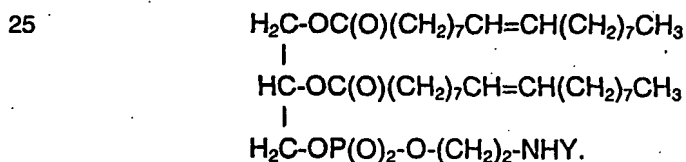
A "blocking group" as used herein refers to a lipid component of the liposome or
 5 cell membrane which sterically or physically blocks a fusogenic component of the membrane from being expressed. Thus, removal of a portion or all of a blocking group by enzymatic activity may also render the liposome or cell membrane fusogenic.

The term "effector" as used in this specification is defined as a compound or
 10 liposome which is added to other liposomes or cells for the purpose of fusing with the other liposome or cell, defined as the "acceptor". The effector liposome thus can be used to deliver a bioactive agent to a cell.

An endocytic vesicle (or endocytic compartment) is a membrane-bound vesicle
 15 that is formed by endocytosis and contains extracellular materials that are to be delivered to any of several destinations within the cell.

An endosome is the cytoplasmic vesicle formed from endocytic vesicles that
 have shed their clathrin coats, or formed directly from coated pits. The pH of
 20 endosomes and endocytic vesicles is acidic.

Preferably, the liposome has a lipid component which comprises a positively charged lipid and a peptide-lipid conjugate having the formula:



More preferably, the peptide comprises the sequence N-methoxy-succinyl-Ala-Ala-Pro-Val and the positively charged lipid is DODAP. Most preferably, presently, the lipid component comprises DODAP and the peptide-lipid conjugate in a respective molar ratio of about 50:50.

Further provided herein is a composition comprising the liposome and a "pharmaceutically acceptable carrier," which is a medium generally acceptable for use in connection with the administration of liposomes to mammals, including humans.

- 5 Pharmaceutically acceptable carriers are formulated according to a number of factors well within the purview of the ordinarily skilled artisan to determine and account for, including without limitation: the particular liposomal bioactive agent used, its concentration, stability and intended bioavailability; the disease, disorder or condition being treated with the liposomal composition; the subject, its age, size and general
- 10 condition; and the composition's intended route of administration, e.g., nasal, oral, ophthalmic, topical, transdermal, vaginal, subcutaneous, intramammary, intraperitoneal, intravenous, or intramuscular (see, for example, Nairn (1985)). Typical pharmaceutically acceptable carriers used in parenteral bioactive agent administration include, for example, D5W, an aqueous solution containing 5% weight by volume of dextrose, and
- 15 physiological saline. Pharmaceutically acceptable carriers can contain additional ingredients, for example those which enhance the stability of the active ingredients included, such as preservatives and anti-oxidants.

- Still further provided is a method of delivering the contents of a liposome to a cell
- 20 which comprises contacting the cell with the liposome of this invention in the presence of a protease capable of cleaving the peptide-lipid conjugate. Delivery can occur *in vitro*, such as for diagnostic purposes or for *ex vivo* delivery of a therapeutic agent or nucleic acid to bone marrow cells. *In vitro* contact of a biological membrane with a lipid-based carrier involves adding the carrier-containing composition of this invention to cultures of
- 25 protease-secreting cells, including various tumor cell lines such as the MCF-7 line, or adding an endogenous protease to the culture medium containing the membranes and the carriers.

- Still further provided is a method of delivering the contents of a liposome to
- 30 nucleated mammalian cells, where at least some of the delivery may occur via an endosomal compartment of the cell. In this method liposomes were developed to take advantage of the low pH of the endosomal compartment in a manner that would

enhance enzyme (elastase) triggering. This is the first report of a liposomal system which can be triggered by physiological levels of elastase to undergo lipid mixing with, and delivery of the liposomes' aqueous contents to nucleated cells. As will be described further in the examples which follow, these activities can occur in two types of human
5 cell lines, a leukemia cell line (HL60) and an adherent cell line, ECV304.

Alternatively, the contacting can be *in vivo*, in which case the cells are preferably mammalian, a pharmaceutically acceptable carrier is used and the liposomes preferably comprise a targeting moiety. *In vivo* administration involves administering the
10 compositions of this invention to the mammal by any of the means, e.g., by intravenous administration, generally accepted in the art for administering pharmaceutical compositions to mammals. The carriers will then circulate in the mammals, and will become fusogenic in the presence of peptidase concentrations sufficient to cleave the carriers' peptide-lipid conjugates; as described hereinabove, such peptidases are found
15 in mammals at, for example, sites of inflammation, microbial infection and tumors. Moreover, incorporation of headgroup-modified lipids into lipid-based carriers increases the amount of time the carriers remain in circulation, and hence the proportion of the administered carrier reaching the intended site of action within the mammal. Furthermore, tumors generally have a higher degree of vasculature than does
20 surrounding tissue, and these blood vessels are typically more permeable to structures such as lipid-based carriers. Accordingly, the carriers accumulate in tumors, thus further enhancing the proportion of administered carrier reaching the intended site of therapeutic action. Fusion *in vivo* can be to the cells secreting the protease as well as to nearby cells in the surrounding tissue.

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In vivo liposomal bioactive agent delivery according to the practice of this invention can deliver therapeutically or diagnostically effective amounts of therapeutic or diagnostic agents into the cells of a mammal afflicted with a disease, disorder or condition amenable to diagnosis or treatment with the agent. Hence, such delivery can
30 be used to diagnose or treat the mammal for the disease, disorder or condition.

The method of this invention can also be used to treat mammals afflicted with inflammatory disorders, by administering to the mammal a liposome containing an anti-inflammation effective amount of an anti-inflammatory agent. Treatable inflammatory disorders include, without limitation, arthritic disorders, autoimmune disorders, 5 atherosclerotic plaque, acute respiratory distress syndrome, inflammatory bowel syndrome, acute nephritis or gout; suitable anti-inflammatory agents include, without limitation, nonsteroidal anti-inflammatory agents, glucocorticoids, bioactive lipids such as ceramides and ether lipids, and prostaglandins. Peptidases known to be present at sites of inflammation include, without limitation: elastase, which recognizes Ala-Ala- and 10 cleaves peptides such as Ala-Ala-, Ala-Ala-Ala-, Ala-Ala-Pro-Val, Ala-Ala-Pro-Met and Ala-Ala-Pro-Ala; stromelysin-1, which recognizes peptides comprising the amino acid sequence Arg-Pro-Lys-Pro-Leu-Ala-Nva-, such as Ac-Arg-Pro-Lys-Pro-Leu-Ala-Nva, MeOSucArg-Pro-Lys-Pro-Leu-Ala-Nva-, carboxy sugar-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Suc-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ac-Ser-Arg-Pro- 15 Lys-Pro-Leu-Ala-Nva-, MeOSuc-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ser-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva-, Ac-Ser-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva- and MeOSuc-Ser-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva, and which cleaves the peptides at the Ala-Nva bond; and, cathepsin G, which is secreted by human neutrophils secreted at the site of inflammation, and cleaves peptides such as Suc-Ala-Ala-Pro-Phe-, carboxy sugar-Ala- 20 Ala-Pro-Phe-, MeOSuc-Ala-Ala-Pro-Met-, Suc-Ala-Ala-Pro-Met-, and carboxy sugar-Ala-Ala-Pro-Met-.

Moreover, peptide substrates for the enzymes granzyme A and granzyme B, secreted by cytotoxic lymphocytes in the synovial fluid of rheumatoid arthritis patients, 25 include, without limitation: Ac-Ala-Ala-Arg-, MeOSuc-Ala-Ala-Arg-, Ala-Ala-Arg-, Ser-Ala-Ala-Arg-, Ac-Ser-Ala-Ala-Arg-, MeOSuc-Ser-Ala-Ala-Arg-, Ser-Ser-Ala-Ala-Arg-, Ac-Ser-Ser-Ala-Ala-Arg-, MeOSuc-Ser-Ser-Ala-Ala-Arg- and carboxyl sugar-Ala-Ala-Arg-, etc. Ac-Ala-Ala-Asp-, MeOSuc-Ala-Ala-Asp-, Ala-Ala-Asp-, Ser-Ala-Ala-Asp-, Ac-Ser-Ala-Ala-Asp-, MeOSuc-Ser-Ala-Ala-Asp-, Ser-Ser-Ala-Ala-Asp-, Ac-Ser-Ser-Ala-Ala-Asp-, 30 MeOSuc-Ser-Ser-Ala-Ala-Asp-, and carboxyl sugar-Ala-Ala-Asp-. Dipeptidylaminopeptidase IV (DAP IV, EC 3.4.14.5), a member of the dipeptidyl peptidase enzyme family, is found in increased concentrations on pig aorta smooth

muscle cells (Palmieri et al., 1989). Vessel wall damage, e.g., after angioplasty or during other inflammatory states exposes the peptidase. For instance, inflammatory edema is associated with breach of the endothelial lining and exposure of smooth muscle cells. Appropriate substrates could be used for liposomal delivery to these sites.

5

The method of this invention can also be used to treat mammals afflicted with cancers, by administering to the mammals a liposome containing an anticancer effective amount of an anticancer agent. Treatable cancers include brain, breast, colon, lung, ovarian, prostate and stomach cancers, as well as sarcomas, carcinomas, leukemias, lymphomas and melanomas; suitable anticancer agents include, without limitation, anthracycline antibiotics, bioactive lipids such as ceramides and ether lipids, taxanes and vinca alkaloids. Peptidases known to be present in the vicinity of tumors include, for example and without limitation: elastase, which cleaves peptides containing the amino acid sequence Ala-Ala-, Ala-Ala-Pro-Val (Nakajima et al., 1979, Castillo et al., 1979); stromelysin-3, which cleaves peptides containing the amino acid sequence Ala-Met; stromelysin-1, which cleaves peptides containing the amino acid sequence Ala-Nva-; human collagenase-3, which cleaves peptides such as MeOSuc-Pro-Cha-Gly-Nva-, Suc-Pro-Cha-Gly-Nva-, Pro-Cha-Gly-Nva-, Pro-Leu-Gly-Leu-, MeOSuc-Pro-Leu-Gly-Leu- and Suc-Pro-Leu-Gly-Leu-; and, the 72-kD gelatinase, which cleaves peptides containing the amino acid sequence Gly-Pro-Gln-Gly-Ile- (see Pei et al., 1994; Knäuper et al., 1996; Boyd, 1996; Unden et al., 1996; and, Kossakowska et al, 1996.) and urokinase plasminogen activator, which cleaves Glu-Gly-Arg and Ac-Lys (Wohl et al., 1980; Johnson et al., 1969; Petkov et al., 1975; Ascenzi et al., 1980), and cathepsin B, which cleaves Arg-Arg (Knight, 1980; Barrett & Kirschke, 1981; Kirschke et al., 1982).

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Moreover, specific peptidases are also found in neuronal tissue (e.g. O'Leary and O'Connor, 1995), suggesting that the liposomes may be designed to treat several neuropathies. Specific aminopeptidases are produced on the membranes of the placental tissue and later secreted suggesting primary localization of this activity in the placenta (Rogi et al., 1996). Several kininases are localized to the kidney. For example renin is found in the zona glomerulosa and/or adrenal medulla (Berka et al., 1996). Certain peptidases have even been identified in skeletal muscle (Ward et al., 1995).

30

Observation of strong activity of an alanylaminopeptidase in the stroma of basal cell carcinoma and DAP IV in the tumor cells themselves (Moehrie et al., 1995) suggest an alanyl-phospholipid or appropriate dipeptides as possible triggers for liposomal fusion
5 with tumor cells.

One of ordinary skill in the art, having chosen a protease from the enzymes described above, or those specifically employed in the Examples described below, would know the cleavage site where cleavage is occurring, and would readily know
10 where the cleavage site is on a particular peptide without undue experimentation.

The method of this invention can also be used to treat mammals afflicted with microbial infections, by administering to the mammals a liposome containing an anti-infection effective amount of an anti-infective agent, such as the various antibiotics. A
15 number of specific peptidases are associated with certain bacteria and may be utilized to deliver liposomal contents to sites of infection (e.g. Spratt et al., 1995). Human immunodeficiency viruses have proteases with particular specificities (e.g. Hoog et al.) that may be expressed in or near infected cells and may be utilized to target fusogenic liposomes for therapy.

20

The contents of the above-cited documents, with their descriptions of secreted enzymes and their target peptides, are incorporated herein by reference.

Liposomal drug delivery according to the practice of this invention can direct the
25 liposomes contents to the vicinity of the target cells. It can also deliver the contents directly into cells, by way of fusion between the liposomes and the cells. "Fusion" of a liposome to a cell involves both binding of the liposome to the cell, as well as mixing of liposomal and cell membrane lipids. Binding and lipid mixing can be assessed by a number of means well known to ordinarily skilled artisans given the teachings of this
30 invention including, for example, those described hereinbelow.

Briefly, liposomes are labeled by incorporation therein of a fluorescent marker and mixed with erythrocyte ghosts, prepared as described hereinbelow. Erythrocyte ghosts are incapable of endocytosis, and hence, any transference of fluorescence between the liposome and ghosts must be due to fusion of the liposome with the plasma membrane. Measurement of erythrocyte ghost fluorescence is thus a measure of the fusion of liposome to the ghosts. Fusion may also occur within the endosome of a living nucleated cell between the liposomal and endosomal membranes, so as to deliver the liposomes' encapsulated material into the cell. Peptidase-mediated cleavage of a peptide-lipid conjugate herein converts a nonfusogenic liposome into a fusogenic liposome. Moreover, the liposome can contain one or more additional fusogenic lipids, including PE's such as DOPE and synthetic lipids such as DOTAP and DODAP. Such lipids promote fusion of their parent liposomes to adjacent lipidic membranes, because of the nonbilayer structures adopted by the lipids in aqueous environments.

However, the peptide-lipid conjugate can also contain a "blocking" group, e.g., a carboxy sugar such as lactobionic acid or N-acetyl neuraminic acid, or a polymeric compound such as a small polyethylene glycol derivative, a polyhydroxyl polymer or a number of other amino acids of a composition containing hydrophilic side chains such as serine or threonine. This blocking group is attached to the N-terminus of the peptide, and inhibits or blocks the liposome and lipidic membrane from approaching closely enough for fusion between the two to occur. Cleavage of the peptide by a protease removes this N-terminal blocking group from the peptide, and hence, allows for fusion between the liposome and the lipidic membrane. Peptidase-mediated cleavage thus, by cleaving the peptide portion of the peptide-lipid conjugate, results in the generation of a fusogenic liposome.

This invention will be better understood in light of the following Examples. However, those of ordinary skill in the art will readily understand that the examples are merely illustrative of the invention as defined in the claims which follow thereafter.

30

EXAMPLES

Example 1

A) Chemical Synthesis of N-Ac-Ala-Ala-DOPE

5 N-acetyl-alanyl-alanyl-dioleoyl phosphatidylethanolamine ("N-Ac-Ala-Ala-DOPE") was synthesized by first preparing an anhydride form of the peptide from N-acetyl-ala-ala-OH, or other suitably blocked carboxyl-terminating peptides; the starting reagent was incubated with N,N-dicyclohexyl carbodiimide (DCC) in the presence of chloroform for a few hours at room temperature. The end-product anhydride is soluble in
10 chloroform, whereas a reaction by-product (dicyclohexyl urea) is not; therefore the anhydride is separated from the undesired by-product by collecting the chloroform and discarding the precipitate. DOPE is added to the anhydride in the presence of triethylamine to catalyze the N-acylation reaction; the mixture is incubated overnight at room temperature. The reaction mixture is applied to a preparative thin layer
15 chromatography (TLC) plate to purify N-acetyl-ala-ala-DOPE, the solvent system being chloroform/methanol/water (65/25/4). The lipid band is identified by spraying the plate with water, after which the band is scraped and solubilized in chloroform/methanol (2/1). Lipid is stored under nitrogen at -70C. Specifically, 25 mg (0.12 mmol) of N-acetyl-alanine-alanine was dissolved in 5 ml of dry tetrahydrofuran (THF), and was stirred for
20 24 hours at room temperature with DCC (25 mg, 0.12 mmol), DOPE (50 mg, 0.26 mmol) and an excess of ethanolamine (ET₃N, 100 ml, 0.7 mmol). The white precipitate of dicyclohexyl urea was filtered through a Celite bed and the filtrate was concentrated. The product was purified by preparative TLC using CHCl₃:MeOH:H₂O (60:15:2) to yield 25 mg as a white flaky powder. The product was characterized by TLC and ³H-
25 NMR (300 MHz, CDCl₃). Purity of the preparation was 90% or greater. The resulting peptide-lipid (N-Ac-Ala-Ala-DOPE) resolved as a single spot by TLC and generated a single peak by reverse-phase HPLC.

B) Chemical Synthesis of 1,2-Dioleoyl-*sn*-glycero-3-phosphoethanolamido-val-pro-ala-ala-sucMeO (MeO-suc-ala-ala-pro-val-DOPE):

30 *i) p-Nitrophenyl-val-pro-ala-ala-sucMeO ester:* To a solution of H-val-pro-ala-ala-sucMeO peptide (540 mg, 1.15 mmol), were added 142 mg (1.38 mmol) of *p*-

nitrophenol, 175 mg (1.38 mmol) of 1,3-dicyclohexylcarbodiimide and a catalytic amount (a few crystals) of 4-dimethylaminopyridine in 10 ml of dry chloroform. The reaction mixture was stirred overnight under nitrogen atmosphere at room temperature. At this point TLC analysis showed that the reaction had gone to completion. The precipitate, 5 dicyclohexylurea (DCU), from the reaction mixture was filtered using a G-2 funnel and the filtrate concentrated under reduced pressure. The residual material was used in the next step without purification; it was characterized by an R_f 0.43 (CHCl_3 :MeOH 9:1 v/v).

ii) 1,2-Dioleoyl-*sn*-glycero-3-phosphoethanolamido-val-pro-ala-ala-sucMeO: To a 10 solution of *p*-nitrophenyl ester of val-pro-ala-ala-sucMeO (600 mg, 1.01 mmol), were added 604 mg (0.81 mmol) of 1,2-dioleoyl-*sn*-glycero-3-phosphoethanolamine and 82 mg (113 ml, 0.81 mmol) of triethylamine in 20 ml of a mixture of the solvents chloroform:tetrahydrofuran (1:4 v/v). The reaction mixture was stirred under nitrogen atmosphere at room temperature overnight. TLC analysis showed that the reaction had 15 gone to completion. The reaction mixture was concentrated under reduced pressure and passed through activated TMD-8 ion exchange resin in THF:H₂O (9:1 v/v).

The phosphorus positive fractions were pooled and concentrated to get the residual product. The residual material was purified by silica gel column 20 chromatography (column was washed with 5% methanol in chloroform, then eluted with CHCl_3 :MeOH:NH₄OH 65:25:4 v/v/v), giving 915 mg (95% yield on the basis of DOPE), which on lyophilization gave a white solid: R_f 0.76 (CHCl_3 :MeOH:NH₄OH 65:25:4 v/v/v) and R_f 0.43 (CHCl_3 :MeOH:H₂O 65:25:4 v/v/v).

25 The lipopeptide molecule gave a positive test for molybdenum reagent and a negative test for ninhydrin reagent. The lipopeptide molecule identity was determined by TLC in two solvent systems: ((i) CHCl_3 :MeOH:NH₄OH 65:25:4 v/v/v and (ii) CHCl_3 :MeOH:H₂O 65:25:4 v/v/v). In both solvent systems the lipopeptide gave a single spot and it is >99% pure. The lipopeptide was characterized by NMR and FAB mass 30 analysis. ¹H-NMR (CDCl_3) some characteristic signals are shown here: δ 0.87 (t, 3H, $J = 7.15$ Hz), 1.27 (40H), 1.56 (4H), 2.0 (8H), 2.23 (t, 4H, $J = 7.15$ Hz), 5.17 (1H), 5.32 (4H, $J = 3.12$ Hz). ³¹P-NMR Spectrum gave single signal. FAB (M^+) calculated for

C62H109N5O15P 1195.55 molecular weight, found mass of 1196.8 (MH+) and mass of 1234.9 (MK+).

C) Preparation of peptide-lipid conjugate: DOPE-Ala-Ala-PEG carboxylate

- 5 A derivative of DOPE was prepared in which a dialanyl peptide linked the DOPE headgroup to a poly(ethyleneglycol) molecule via amide linkages at each end of the peptide. This is referred to a "DOPE-ala-ala-PEG-carboxylate". For the synthesis, poly(ethyleneglycol) bis(carboxymethyl) (Aldrich) was utilized, a polyethylene glycol with carboxyl groups at each terminus and a molecular weight of 228. To a solution of
- 10 poly(ethyleneglycol) bis(carboxymethyl) ether (8.5 mg, 0.034 mmole) in 2 ml of chloroform, was added dicyclohexyl carbodiimide (DCC, 5.0 mg, 0.023 mmole). To this solution the peptide-lipid, N-ala-ala- 1,2-dioleoyl-sn-glycero-3-phosphatidylethanolamine (20 mg, 0.023 mmol), was added in chloroform to give a final volume of 3.0 ml. The reaction mixture was stirred overnight under nitrogen atmosphere at room temperature.
- 15 At this point thin layer chromatography (TLC) was performed. The precipitate, DCU, from the reaction mixture was filtered using a G-2 funnel and the filtrate concentrated under reduced pressure. TLC indicated the necessity to further purify the preparation. The remaining material was further purified by silica gel column chromatography, as in section B of this Example, to remove residual starting material and yielded 6.5 mg of
- 20 purified material (>99%) with an R_f of approximately 0.28 on a silica gel TLC plate run in the solvent CHCl₃/methanol/NH₄OH (65/25/4). As expected this purified product was more polar than the starting peptide-lipid. By TLC this product gave a single spot and >99% purity. The product was also characterized FAB mass analysis. The calculated FAB (M+) was 996 and a mass of 1134.6 (MK+) was observed.

25

Example 2

Elastase and proteinase K-mediated cleavage of N-Ac-ala-ala-DOPE

- Cleavage of N-Ac-ala-ala-DOPE to DOPE by elastase was monitored by thin layer chromatography (TLC). The chemical structure of N-Ac-ala-ala-DOPE and its
- 30 postulated scheme of conversion to DOPE by enzymatic cleavage is shown in Fig. 1. 100-200 nmol of N-Ac-AA-DOPE SUVs were incubated with 1 mg enzyme in 0.1 ml overnight at 37°C. Human leukocyte elastase was purchased from Calbiochem (San

Diego, CA). Lipid was extracted by organic phase separation (Bligh and Dyer, 1959) twice. Collected lipid was dried under N₂ stream and exposed to vacuum for 4 hours-overnight. Samples were resuspended in chloroform and spotted onto TLC plates. TLC was run using chloroform/methanol/water (65:25:4), air dried, sprayed with molybdenate
5 blue, and charred on a hot plate. Treatment of N-Ac-AA-DOPE liposomes with elastase generated a product corresponding to DOPE, whereas untreated N-Ac-AA-DOPE showed no change (Fig. 2A). Therefore elastase recognized N-Ac-AA-DOPE and cleaved the dipeptide to yield DOPE.

10 Several proteases were tested to determine whether an enzyme with similar substrate specificity could be used as a model for elastase mediated cleavage of N-Ac-AA-DOPE. Proteinase K is a serine protease that, similarly to elastase, can cleave at peptide bonds C-terminal to aliphatic residues. Upon incubation of N-Ac-AA-DOPE liposomes with proteinase K the peptide-lipid was cleaved and DOPE was generated
15 (Fig. 2B).

Subsequently, proteinase K, which has a similar substrate specificity to elastase, was used to model the elastase effect because of the lower cost of the enzyme. Referring to Fig. 3, samples A and C contained pure N-acetyl-ala-ala-DOPE and pure DOPE, respectively; sample B
20 consisted of multilamellar liposomes, prepared as described hereinbelow, composed of pure N-acetyl-ala-ala-DOPE (2 μ mole total phospholipid), which were incubated in 200 μ l of 154 mM NaCl, 10 mM TES, 0.1 mM EDTA at pH 7.4 and 37 °C for two hours in the presence of 1 mg of proteinase K. After the first incubation, another equal amount of proteinase K was added and the sample was incubated for another 2 hours. All three samples were dissolved into a final
25 concentration of 5% deoxycholate, 50 mM EDTA and 10 mM HEPES at pH 8 for proton decoupled ³¹P NMR. Integration showed that approximately 20% of the N-acetyl-ala-ala-DOPE was converted to pure DOPE in sample B.

The conversion of N-Ac-AA-DOPE to DOPE was also monitored by ³¹P-NMR
30 analysis. N-Ac-AA-DOPE LUVs were prepared and treated with or without proteinase K (1.5 mg protein/100 nmol lipid) overnight at 37°C. Samples were mixed with buffer (10% deoxycholate, 100mM EDTA, 20mM Hepes) and deuterium oxide (Cambridge Isotope Laboratories, Woburn, MA) (1:4:2) and transferred to 5mm NMR tubes. Samples were

monitored at room temperature in a Bruker AC300 spectrometer operating at 121.5 MHz, with 110ms 90° radio frequency pulse for proton decoupling and set to 2 sec interpulse delay to avoid signal saturation. Sweep width was set at 50kHz. 1 Hz line broadening was applied to all spectra. N-Ac-AA-DOPE liposomes treated with
5 proteinase K (1.5 mg protease/100 nmol lipid) resulted in the appearance of a peak 0.3 ppm upfield from N-Ac-AA-DOPE, corresponding with pure DOPE.

Elastase and proteinase K mediated cleavage of N-Ac-AA-DOPE was quantitated using liposomes composed of N-Ac-AA-DOPE and DOTAP, a positively
10 charged lipid. DOTAP was included to provide a counterbalancing positive charge, and was used as a standard by which different samples could be normalized and compared. After treatment with elastase or proteinase K the reduction in the amount of N-Ac-AA-DOPE was monitored by HPLC (Fig. 4). Liposomes composed of DOTAP/N-Ac-AA-DOPE (1:1) were incubated with enzyme under given conditions. Lipid was extracted by
15 the Bligh-Dyer procedure twice.

Collected lipid was dried under a N₂ stream and exposed to vacuum for 4 hours-overnight. Samples were resuspended in 100% ethanol and injected in 30 ul aliquots into Spherisorb silica columns (150 x 4.6mm, 0.3 um, Keystone Scientific). HPLC was
20 performed using a hexane:isopropanol:water:TFA mobile phase. Hexane and TFA were held constant at 37% and 0.2%, respectively. The N-Ac-AA-DOPE peak was detected using a gradient of 59-55% isopropanol:4-8% water. Flow rate was 1.5 ml/min, column temperature was set at 45°C, and peaks were detected by a UV detector set at 205nm. Lipid peaks were quantitated in comparison to standard curves generated by injecting 5-
25 200 nmol of DOTAP or N-Ac-AA-DOPE and monitoring 205nm signal. Percent cleavage was calculated by normalizing peaks to DOTAP, then determining the decrease in N-Ac-AA-DOPE peak size relative to starting amounts.

Both elastase and proteinase K cleaved N-Ac-AA-DOPE to a similar extent (Fig.
30 4). To verify that the cleavage of N-Ac-AA-DOPE was due to proteinase K enzymatic activity, liposomes were treated with heat inactivated proteinase K. Proteinase K was inactivated by heating at 95°C for 1 hour, after which the enzyme was incapable of

cleaving the chromogenic substrate *N*-Ac-AAA-pNA. Treatment of DOTAP/*N*-Ac-AA-DOPE liposomes with heat inactivated proteinase K did not result in any cleavage of *N*-Ac-AA-DOPE (Fig. 4), indicating the requirement for active proteinase K. Since proteinase K has been shown to share substrate specificity with elastase and is considerably less costly than human leukocyte elastase, several experiments were conducted with proteinase K. Later experiments on the effects of human leukocyte elastase on *N*-Ac-ala-ala-DOPE used thin layer chromatography ("TLC"), a high sensitivity method, to detect the product of enzyme activity. In these experiments liposomes, prepared as described hereinbelow, were composed of 100% *N*-Acetyl-ala-ala-DOPE and were then extruded to approximately 0.2 microns diameter/ Fifty micrograms of liposomal lipid was incubated alone or with one unit of HLE overnight at 37°C.

Example 3

15 **Cleavage of MeO-suc-ala-ala-pro-val-DOPE by human leukocyte elastase (HLE)**

A) HLE dose titration

To determine if the peptide-lipid MeO-suc-ala-ala-pro-val-DOPE is also a suitable substrate for elastase mediated cleavage 50 nmol of MeO-suc-ala-ala-pro-val-DOPE liposomes (SUVs) were incubated with 0, 2.5, 5, 10, or 20ug HLE (from Calbiochem; 20 units/mg protein; 1 unit = amount of enzyme that will hydrolyze 1.0 umol of MeO-suc-ala-ala-pro-val-pNA per min at 25 degrees C, pH 8.0) overnight at 37 degrees C in 50 ul volume of 10mM TES/ 154 mM NaCl/ 0.1mM EDTA, pH 7.4, containing 1.5mM Ca and 1.5 mM Mg.

25 Lipid was extracted using the Bligh-Dyer technique (chloroform/methanol/water : 2/1.7/1), dried under nitrogen, placed under high vacuum for ~ 3 hours. Samples were resuspended in 5 ul chloroform and spotted onto TLC plates. 20ug of pure DOPE was also spotted for comparison purposes. TLC solvent system was chloroform/methanol/ammonium hydroxide (65/25/5). Plates were air dried, sprayed with molybdenate blue, then charred at 180 degrees C.

As shown in Fig. 5A, between 5 -40 micrograms of HLE were effective in cleaving the AAPV-PE (compare lanes 2,3,5 & 6 with lane 4 & 8, and with the proteinase K control, lane 7).

- 5 **B) Kinetics of HLE-mediated cleavage of MeO-suc-ala-ala-pro-val-DOPE:** MeO-suc-ala-ala-pro-val-DOPE liposomes were incubated with 0 or 5 ug HLE for 1, 2, 4 hours, or overnight and processed as above. The peptide was cleaved by HLE in as little as 1 hour at 37 degrees C (Fig. 5B, lane 2), suggesting that the cleavage of MeO-suc-ala-ala-pro-val-DOPE occurs within a physiologically relevant time frame.

10

C) Cleavage of MeO-suc-ala-ala-pro-val-DOPE by human neutrophil granule proteins

- Since elastase is produced by activated neutrophils the cleavage of MeO-suc-ala-ala-pro-val-DOPE by unpurified granule proteins was monitored to mimic more closely the in vivo situation. Neutrophils were obtained from human whole blood by standard procedures employing density centrifugation. Granules were isolated from these neutrophils by centrifugation following nitrogen cavitation of cells, again following established procedures. Protein concentration of neutrophil granules was determined after repeated freeze-thawing of granules to release proteases.

20

- Fifty nmol MeO-suc-ala-ala-pro-val-DOPE liposomes (SUVs) were incubated with 0, 2.5, 5, 10, or 20ug neutrophil granule proteins overnight at 37 degrees C in 50 ul volume of 10mM TES/ 154 mM NaCl/ 0.1mM EDTA, pH 7.4, containing 1.5mM Ca and 1.5 mM Mg. Samples were processed as described above. The results shown in Fig. 5C indicate that 2.5 µg of neutrophil granule proteins were sufficient to detect cleavage of MeO-suc-ala-ala-pro-val-DOPE to DOPE, suggesting that crude neutrophil granule proteins can convert the peptide-lipid to DOPE, and therefore liposomes containing this peptide-lipid can be activated to fuse under physiological conditions.

30

- The conversion of MeO-suc-AAPV-DOPE to DOPE was also quantitated by ³¹P-NMR. Since subsequent experiments employed liposomes containing both DODAP and MeO-suc-AAPV-DOPE, vesicles prepared at a 1:1 (mol:mol) ratio of these two

components were prepared by freeze-thaw/extrusion method as described above, and incubated at 37°C for 2 hours with increasing amounts of elastase (0 micrograms to 5 micrograms elastase/100 nmol lipid). ³¹P-NMR analysis demonstrated an elastase concentration dependent cleavage of MeO-suc-AAPV-DOPE and appearance of DOPE (Fig. 6, solid line). A small shoulder that may indicate an incomplete peptide cleavage product was also observed near the original peptide-lipid peak. Treatment with 5 micrograms of elastase/100 nmol lipid yielded 20% DOPE. Longer incubation may have lead to further digestion, though multiple lamellae and/or the surface charge of the liposome may limit the ultimate amount digested. The maximum exposed peptide-lipid for intact liposomes would occur with unilamellar vesicles. Assuming only the outer leaflet peptide-lipid in unilamellar vesicles is available for digestion, the minimal percentage conversion of exposed MeO-suc-AAPV-DOPE to DOPE in the outer leaflet lipid of this preparation was 40% (Fig. 6, dotted line). If the average number of lamellae were greater than one, the percentage conversion was even higher. The ratio of encapsulated volume to total lipid would appear to indicate an average of approximately 2.5 lamellae per vesicle for this preparation which would indicate that 100% of available peptide-lipid had been cleaved under these conditions. The concentration of elastase to produce this amount of cleavage (12.5 micrograms elastase/ml) is less than the effective concentration of elastase activity found in the epithelial lining fluid from patients with cystic fibrosis (Birrer et al, 1994). The contents from human neutrophil granules also cleaved MeO-suc-AAPV-DOPE and generated DOPE. The number of neutrophils required to observe this effect was less than that observed in epithelial lining fluid from cystic fibrosis patients (Birrer et al., 1994), indicating the amount of elastase required to cleave MeO-suc-AAPV-DOPE is within concentrations that are physiologically or therapeutically relevant

25

Example 4

Enzyme-mediated cleavage of N-Acetyl-ala-ala-PE: To compare the effect of different proteolytic enzymes on the cleavage of N-acetyl-ala-ala-DOPE, small unilamellar liposomes composed of DOTAP and N-acetyl-ala-ala-PE (1:1 molar ratio, 100 nmole), prepared as described hereinabove, were incubated overnight at 37 degrees C. with either 0.5 mg (experiment 1) or 1.0 mg (experiment 2) of: S. caespitosus protease S.

30

griseus ("pronase"), peptidase, chymotrypsin, trypsin, protease type I, proteinase K, or with no enzyme added.

Subsequently, lipid was extracted from these preparations by organic phase separation, and dried under vacuum. The dried lipid was suspended in ethanol, and the resulting lipidic solution was then injected into a normal phase silica column. Analysis was by HPLC, as described above. Lipid peaks were quantitated, and the amount of lipid cleavage was calculated as a percentage of the starting lipid. Fig. 7 shows that both proteinase K, with a known cleavage site, and the nonspecific mixture of proteases found in pronase, effected cleavage of N-Ac-ala-ala-DOPE from the liposomes, and the greater release being obtained with the increased enzyme concentrations. Other proteases, however, did not cleave the N-Ac-ala-ala-DOPE from the liposomes.

Example 5

15 - Liposome Preparation

NBD/Rh labeled or unlabeled large unilamellar vesicles (LUVs) were prepared as described before (Mayer et al.). Briefly, the lipid mixture in chloroform was dried under a nitrogen stream to a thin film, which was then left under vacuum overnight to remove residual solvent. The lipid film was hydrated with TES buffered saline (10 mM TES, 0.1 mM EDTA, 154 mM NaCl, pH 7.4). Brief vortexing was applied to ensure complete hydration. After ten cycles of freeze/thaw in liquid nitrogen/room temperature water bath, the sample was extruded ten times through 0.1 μ m polycarbonate membrane filter (Poretics Corp., Livermore, CA). The liposomes were stored at 4°C. Multilamellar vesicles were prepared by hydrating the dried lipid film.

25

The phospholipid concentration of each liposome preparation was determined by phosphate assay (Bartlett). The approximately 0.1 μ m size of the liposomes was confirmed on a Nicomp submicron particle sizer (Nicomp Instruments, Inc., Goleta, CA) using quasi-elastic light scattering.

30

Example 6

Preparation of Resealed and Unsealed Human Erythrocyte Ghosts

Resealed ghosts are referred to as erythrocyte ghosts unless otherwise specified, and were prepared as previously described (see Williamson et al.; Clague et al., the contents of which are incorporated herein by reference). Briefly, fresh human blood was washed several times with cold 10 mM TES buffered saline to remove plasma and white cells. Then 2 ml of washed erythrocytes (50% hematocrit) were pre-swelled in cold hypotonic solution containing 8 ml H₂O and 9.6 ml 10 mM TES buffered saline, and then pelleted at 850xg for 5 minutes. The pellet was resuspended in 40 ml cold lysis buffer (10 mM Tris, 0.1% bovine serum albumen (BSA), 2 mM MgCl₂, and 0.1 mM EGTA) and incubated on ice for at least 2 minutes. After addition of 4.5 ml 10x resealing buffer (1.22 M NaCl, 30 mM KCl, 0.15 M Na₂HPO₄, 50 mM KH₂PO₄, and 2 mM MgCl₂), the sample was incubated at 37°C for 40 minutes. The resealed ghosts were pelleted at 1750xg for 10 minutes and washed several times until no hemoglobin could be observed in the supernatant. The ghosts were stored at 4°C and used within one week. A prothrombinase assay was used to test for maintenance of phospholipid asymmetry in the ghosts resulting from this method of preparation, as described below.

Example 7**Prothrombinase Assay**

The phospholipid asymmetry of the erythrocyte membranes was measured by a
5 prothrombinase assay, which detects the presence of PS (phosphoserine) in the outer monolayer
of the membrane. The assay was performed as previously described (Wilson et al., 1993, the
contents of which are incorporated herein by reference) with some modifications. Briefly, 4×10^{15}
erythrocyte ghosts were incubated at 37°C for 3 minutes in Tris buffer (50 mM Tris-HCl, 120 mM
NaCl, pH 7.4) containing 6 mM CaCl_2 , 0.33 unit ml^{-1} factor V/Va, 0.33 unit ml^{-1} factor Xa, and 1.3
10 unit ml^{-1} prothrombinase; the total reaction volume was 1 ml. After 15 mM EDTA was added to
stop the reaction, the cells were pelleted by centrifugation. About 900 μl supernatant was mixed
with 100 μl chromogenic substrate, sarcosine-Pro-Arg-*p*-nitroanilide (500 μM) and OD_{405} was
measured kinetically. The rates ($\Delta\text{OD}/\text{min}$) of intact erythrocytes and unsealed erythrocyte ghosts
were taken as 0% and 100% accessible PS, respectively. Using this scale, the accessible PS on
15 the resealed ghosts was found to be about 16%.

Example 8**Design of fusion-triggerable liposomes containing *N*-Ac-AA-DOPE**

The threshold of fusogenicity was determined by preparing liposomes with
20 increasing amounts of PE transesterified from egg PC. This PE was preferred over
DOPE because of its higher H_{II} transition temperature ($\sim 37^\circ\text{C}$ vs. 10°C , respectively),
which aids in the preparation of stable liposomes yet does not inhibit fusion. DOTAP
was chosen as the positively charged lipid. Fusion assays were performed for
DOTAP/*N*-Ac-AA-DOPE/PE liposomes containing the fluorescent membrane probes *N*-
25 NBD-PE ((*N*-(7-nitrobenz-2-oxa-1,3-diazol-4-yl) PE) and *N*-Rho-PE (*N*-
lissamineorhodamine B sulfonyl (PE)) and inversely varying amounts of *N*-Ac-AA-DOPE
and PE. These liposomes were monitored for lipid mixing with either unlabeled target
liposomes or for lipid mixing and binding with RBC ghosts. Lipid mixing between
NBD/Rh labeled liposomes and unlabeled ghosts was measured in 10 mM TES buffered
30 saline by the NBD/Rh resonance energy transfer (RET) assay (Struck et al., the
contents of which are incorporated herein by reference).

Liposomes were prepared with 1 mol% N-NBD-PE and 1 mol% N-Rho-PE, which results in quenching of the N-NBD-PE fluorescence signal. Membrane fusion results in probe diffusion and relief from self-quenching, which is monitored as an increase in N-NBD-PE fluorescence. Liposome-liposome lipid mixing was initiated by addition of 10 nmol of fluorescently labeled liposomes to 90 nmol unlabeled liposomes in microcentrifuge tubes containing 1 ml of TES/NaCl/EDTA buffer with 1.5mM Ca⁺⁺/1.5mM Mg⁺⁺. For fusion with cells 1x10⁸ RBC ghosts were substituted for unlabeled liposomes. All samples were shaken in an Eppendorf Thermomixer (Brinkmann Instruments, Inc., Westbury, NY), 700 rpm/min, during the 37°C incubation for 30min. N-NBD-PE fluorescence was monitored in a T-format PTI Alphascan spectrofluorometer (Princeton, NJ) with a xenon short arc lamp using 450 nm excitation/530nm emission wavelengths and 5nm slitwidths. 450 nm band pass and 500nm cutoff filters were utilized for excitation and emission light paths, respectively, to reduce stray light. Maximal fluorescence dequenching was determined by addition of 0.1% C12E8 detergent (octaethylene glycol monododecyl ether).

The threshold for fusogenicity depends upon the target in question. Liposomes composed of DOTAP (20%) and PE (80%) fused with both target liposomes and RBC ghosts. Inclusion of 10 mol% N-Ac-AA-DOPE with a corresponding decrease in PE to 70 mol% generated liposomes that were still capable of fusing with PE/PS liposomes but not PC/PS liposomes (Fig. 8A). The requirements for membrane fusion with RBC ghosts appeared to be more stringent, with inclusion of 5 mol% N-Ac-AA-DOPE inhibiting both the lipid mixing and the binding significantly (Fig. 8B). Defining the different threshold of fusion for different targets creates a gradient of sensitivity for fusion that can be used to determine optimum conditions for activating N-Ac-AA-DOPE containing liposomes to fuse. As PE/PS liposomes appeared to be the most sensitive target, we focused on a composition of DOTAP/N-Ac-AA-DOPE/PE liposomes that could be activated to fuse. The threshold of PE content appeared to be between 65-70 mol%. In order to create a liposome that is not initially highly positively charged, DOTAP and N-Ac-AA-DOPE were added in equivalent amounts to yield liposomes composed of DOTAP/N-Ac-AA-DOPE/PE in a 15/15/70 mol ratio.

These results suggests that the threshold PE content is between 65-70 mol% for fusion with PS/PE liposomes, but 70-80% for fusion with erythrocyte ghosts.

Example 9

5 **Concentration-dependence and time-dependence of proteinase K activity**

Having established the liposome compositions close to the threshold for fusion, the concentration and time-dependence of proteinase K activation of fusion was determined. Large unilamellar liposomes were prepared and fluorescently labeled, as described hereinabove, so as to contain DOTAP, N-acetyl-ala-ala-PE and tPE

10 ("transphosphatidylated PE," PE prepared from egg PC by transphosphatidylation reaction; i.e., phospholipase D, treatment of egg PC in presence of phosphoethanolamine) in a 15:15:70 mole % ratio. For fusion experiments, 1 mole% of N-NBD-PE and 1 mole % of N-Rho-PE were added as fluorescence probes. For cleavage experiments, liposomes without the fluorescent probes were used.

15

Aliquots of these liposomes were then incubated at 37 degrees C. overnight with: 0, 0.1, 0.25, 0.5 or 1.0 mg of proteinase K (100 nmoles lipid/1mg protein).

Subsequently, aliquots of these effector liposome preparations were incubated with unlabeled PE/PS acceptor liposomes, at a ratio of 1:10 effector liposomes:acceptor
20 liposomes. The degree of lipid mixing between effector and acceptor liposomes, assessed as described hereinabove, was then determined as a measure of interliposome fusion

Large unilamellar liposomes were prepared as described hereinabove to contain
25 DOTAP and N-acetyl-ala-ala-PE in a 1:1 molar ratio. Lipid extraction and HPLC analysis was then performed, as described hereinabove, to assess the extent of lipid cleavage. The reliance of fusion activation upon enzyme cleavage of N-Ac-AA-DOPE was further assessed by examining the concentration and time dependencies of both events. DOTAP/N-Ac-AA-DOPE/PE liposomes were either incubated with 0, 0.1, 0.25,
30 0.5, and 1 mg proteinase K/100nmol lipid overnight, or with 1 mg proteinase K/100nmol lipid for 1, 2, 4 hours or overnight. These liposomes were monitored for N-Ac-AA-DOPE cleavage by HPLC or for lipid mixing with acceptor liposomes by N-NBD-PE

fluorescence dequenching. A similar concentration dependence was evident for both *N*-Ac-AA-DOPE cleavage and liposome fusion (Fig. 0A). Treatment with 0.5 or 1 mg proteinase K yielded apparently maximal cleavage and fusion activity. Only background levels of both activities were observed when 0 or 0.1 mg of the enzyme were used. The kinetics of proteinase K mediated cleavage and fusion activation were also correlated, with overnight incubation giving the highest amount of cleavage and lipid mixing (Fig. 9B, and data described in Example 2). These results further support the contention that the activation of fusion of DOTAP/*N*-Ac-AA-DOPE/PE liposomes is due to enzymatic cleavage of *N*-Ac-AA-DOPE.

10

As shown in Fig. 8A, between 0.5 - 1.0 mg of proteinase K effected maximal cleavage and fusion activity, and that lipid mixing is a good indicator for monitoring *N*-Ac-AA-DOPE cleavage. Fig. 8B shows no significant differences when liposomes are incubated with proteinase K for a period of between 1 and 4 hours, and that maximal release and fusion activity, as measured by both methods, is maximal after an overnight incubation.

Example 10

20 **Activation of liposome-liposome lipid mixing by elastase and proteinase K cleavage**

Since elastase and proteinase K were capable of cleaving *N*-Ac-AA-DOPE to DOPE (Figs. 2 & 4) both enzymes were tested for their ability to activate DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes to fuse. These liposomes were treated overnight at 37°C with elastase, or proteinase K, or without either enzyme, after which liposomes were incubated with PE/PS liposomes and lipid mixing monitored by relief of *N*-NBD-PE fluorescence quenching. Liposomes were incubated with protease at a 1 mg protease/100 nmol lipid/0.1 ml buffer ratio, unless otherwise stated. This concentration of proteinase K was found to have comparable activity, within an order of magnitude, with that of elastase in rheumatoid arthritis synovial fluid (Al-Haik et al., 1984, Agents Actions 15:436-442; and our unpublished data). Mixtures were incubated at 37°C in microcentrifuge tubes with constant shaking in an Eppendorf Thermomixer, 700

30

rpm/min. Treated liposomes were then assayed for *N*-Ac-AA-DOPE cleavage by HPLC, as described above. For fusion experiments liposomes containing fluorescent membrane probes were treated with protease and then the concentrations of liposomes were determined by monitoring direct *N*-Rho-PE fluorescence (550ex/590em) and
5 comparing with a known amount of stock liposomes. Aliquots of these fluorescently labeled protease treated liposomes were incubated with unlabeled target liposomes or cells and lipid mixing was determined as described above.

Treatment by either enzyme resulted in a greater extent of lipid mixing over that
10 of untreated liposomes (Fig. 10). This result, coupled with the shared substrate specificity of proteinase K and elastase, suggests proteinase K activation serves as a suitable substitute for elastase to characterize the fusion activation of *N*-Ac-AA-DOPE containing liposomes.

15 A causal relationship between cleavage of the *N*-Ac-AA-DOPE peptide-lipid and fusion activation of DOTAP/*N*-Ac-AA-DOPE/PE liposomes was studied using heat inactivated proteinase K. DOTAP/*N*-Ac-AA-DOPE/PE (15/15/70 mol%) liposomes containing the fluorescent membrane probes *N*-NBD-PE and *N*-Rho-PE were incubated overnight at 37°C with active or heat inactivated proteinase K, after which an aliquot of
20 the liposomes was incubated with unlabeled PS/PE acceptor liposomes to monitor the extent of lipid mixing. Treatment of DOTAP/*N*-Ac-AA-DOPE/PE liposomes with active proteinase K resulted in ~30% fluorescence dequenching, a six-fold increase in lipid mixing over the untreated liposomes (Fig. 11). However, treatment with an identical amount of the heat inactivated enzyme did not activate liposomes to fuse (Fig. 11).
25 Therefore, enzymatic activity is essential for the liposomes to become fusogenic, indicating *N*-Ac-AA-DOPE cleavage is crucial for triggering the fusogenic potential.

Example 11

Activation of DOTAP/*N*-Ac-AA-DOPE/PE fusion with RBC ghosts

30 Since DOTAP/*N*-Ac-AA-DOPE/PE liposomes could be activated to fuse with target liposomes after enzymatic cleavage, we determined if activated fusion of *N*-Ac-AA-DOPE containing liposomes with cell membranes could also be observed. As fusion

with RBC ghosts (Fig. 8B) appeared to exhibit a different threshold of fusogenicity than liposomes (Fig. 8A), we prepared DOTAP/*N*-Ac-AA-DOPE/PE liposomes at a 20/10/70 mol ratio. The overall positive charge of these liposomes improves the binding to cell membranes, relative to the 15/15/70 composition, without permitting the liposomes to fuse with cell membranes in the absence of an activating trigger (Fig. 8B). After an overnight, 37°C, incubation of these liposomes with proteinase K, lipid mixing with RBC ghosts was observed in the presence of the protease inhibitor phenyl methyl sulfonyl fluoride (PMSF) (Fig. 12, with Proteinase K). The activity of residual proteinase K transferred from the initial incubation was negligible (Fig. 12, Proteinase K control).

Specific activation of DOTAP/*N*-Ac-AA-DOPE/PE (20/10/70 mol%) fusion with RBC ghosts was also observed under continuous kinetics conditions. Only liposomes pretreated with proteinase K were capable of fusing with RBC ghosts while untreated liposomes did not (Fig. 13). The addition of active proteinase K to untreated liposomes also did not induce fluorescence dequenching (Fig. 13, curve c), indicating the observed increase for proteinase K treated DOTAP/*N*-Ac-AA-DOPE/PE liposomes was due to specific fusion activation.

To determine if the lipid mixing observed after proteinase K activation was due to true fusion of liposomes with cell membranes and not potential artifacts of the lipid mixing assay, such as membrane probe exchange or hemifusion between outer leaflet membranes, DOTAP/*N*-Ac-AA-DOPE/PE (20/10/70 mol%) liposomes were loaded with 10,000 MW fluorescent aqueous probe TX-red dextran. Liposomes were then treated with proteinase K and incubated with RBC ghosts. After washing extensively to remove unbound liposomes the RBC ghosts were observed under fluorescence microscopy. The pellet was resuspended in 0.1 ml buffer and observed under an Olympus BH-2 fluorescence microscope (Olympus Corp., Lake Success, NY) using an apochromat 40x oil (N.A. 1.00) objective. TX-red fluorescence was excited by a xenon lamp transmitted through a green excitation cube (580nm dichroic mirror, 545nm excitation filter). Non-fluorescent images were observed with transmitted light Nomarski differential interference contrast microscopy.

Bright diffuse fluorescence could be observed in a portion of the ghosts (Fig. 14), indicating complete fusion occurred between liposomes and certain ghosts with subsequent transfer of the fluorescent aqueous probe. Differences in fluorescence levels may be due to differences in the number of liposomes fusing with a single ghost.

- 5 The observed fluorescence does not appear to be due to non-specific uptake of dextran out of leaky liposomes, as incubation of RBC ghosts with unlabeled liposomes and free TX-red dextran did not result in observable aqueous probe labeling (Fig. 14). Thus DOTAP/N-Ac-AA-DOPE/PE liposomes can be activated by enzymatic cleavage of the peptide-lipid to fuse with cell membranes and deliver their aqueous contents.

10

Example 12

Cells

- HL60 human leukemia and ECV304 human cells were obtained from the American Type Culture Collection (ATCC, Rockville, MD). HL60 cells were passaged as
15 suspension cultures in RPMI 1640 medium supplemented with 10% heat-inactivated fetal bovine serum (FBS). Adherent ECV304 cells were grown in Medium 199 supplemented with 10% heat-inactivated FBS. Greater than 98% viability was observed during routine tissue culture. RPMI 1640, FBS and Hanks' Balanced Salt Solution (HBSS) were purchased from Life Technologies, (Gaithersburg, MD).

20

Example 13

Activation of MeO-suc-ala-ala-pro-val-DOPE containing liposomes by HLE for enhanced binding/fusion to HL60 cells

- 100 nmol of DOTAP/ AAPV-PE (3:1; 1:1; or 1:3) were incubated with or without
25 HLE (5ug/100nmol lipid) in 100 ul for 2 hours at 37C, pH 7.4 with constant shaking. Afterwards 10nmol of liposomes with or without HLE pretreatment were incubated with $1-2 \times 10^6$ HL60 (human leukemia) cells at pH 7.4 or 4. Samples were incubated 2 hours at 37C with constant shaking. After incubation, cells were washed once, resuspended in 0.5 ml, and transferred to wells of Falcon 24 well plate (PRIMARIA [Trademark],
30 Becton-Dickinson Co., Lincoln Park, NJ). Fluorescence was monitored with or without detergent in a Cytofluor plate reader (Perseptive Biosystems, Framingham, MA).

Liposome input (no cells) were not washed and were transferred directly to the wells of 24 well plates.

The results of a series of experiments are described below in Table 1 In a first series of experiments, liposomes exhibited higher binding at pH 4.0. Binding also appeared to be higher after HLE pretreatment. In addition, DODAP/AAPV-PE (1:1) may possibly exhibit higher fluorescence dequenching after HLE treatment, suggesting lipid mixing between liposomes and cells had occurred.

<u>DODAP/AAPV-PE</u>	<u>% lipo bound</u>		<u>% NBD</u>	
	<u>pH 7.4</u>	<u>pH 4.0</u>	<u>FDQ</u> <u>pH 7.4</u>	<u>pH 4.0</u>
1:3 - HLE	2.68	53.58	104.22	33.13
1:3 + HLE	0.74	70.34	-32.43	32.53
1:1 - HLE	2.37	56.95	59.94	23.86
1:1 + HLE	2.87	85.91	77.42	44.81
3:1 - HLE	6.76	67.88	-5.05	64.52
3:1 + HLE	7.78	80.82	43.43	69.04

10

Three separate experiments also displayed enhanced binding of DODAP/AAPV-PE (1:1) to HL60 cells. One showed an enhancement from 64% to 86% fluorescence dequenching. In terms of total number of liposomes fused, HLE pretreatment appeared to enhance lipid mixing in all three experiments.

	<u>w/o HLE</u>	<u>w/ HLE</u>
% bound	20.29	45.22
# bound	1.22E+10	2.72E+10
% lipid mixing	48.06	44.24
# lipid mixed	5.87E+11	1.20E+12

	<u>w/o HLE</u>	<u>w/ HLE</u>
% bound	7.72	33.17
# lipos bound	4.60E+09	2.00E+10
% lipid mixing	57.12	25.89
# lipid mixing	2.65E+11	5.17E+11

	<u>w/o HLE</u>	<u>w/ HLE</u>
% bound	38.09	46.89
# Lipos bound	2.29E+10	3.08E+10
% lipid mixing	63.74	86.25
# lipid mixed	1.46E+12	2.43E+12

5 Example 14.

Binding and lipid mixing of liposomes to HL60 cells

In another series of experiments, lipid mixing was monitored by the *N*-NBD-PE/*N*-Rho-PE resonance energy transfer assay, as described [Struck et al., 1981].

- Liposomes were prepared with 0.75 mol% *N*-NBD-PE and 0.75 mol% *N*-Rho-PE, which
 10 results in quenching of the *N*-NBD-PE fluorescence signal. Membrane fusion results in probe diffusion and relief from self-quenching, which is monitored as an increase in *N*-NBD-PE fluorescence. DODAP/MeO-suc-AAPV-DOPE (1:1 mol:mol) liposomes were incubated in TES/NaCl/EDTA buffer with or without elastase (5 • g/100 nmol lipid, 250 • M lipid concentration) for 2 hours at 37°C, pH 7.4. HL60 cells were washed with
 15 TES/NaCl/EDTA buffer and incubated with liposomes (1x10⁶ cells, 10 nmol liposome) in 200 • l TES/NaCl/EDTA buffer. Samples were either at pH 7.4 or adjusted to pH 5 by

the addition of dilute HCl. All samples were shaken in an Eppendorf Thermomixer (Brinkmann Instruments, Inc., Westbury, NY), 700 rpm/min, for 30 min at 37°C. There was no reduction in cell viability following this procedure, as detected by trypan blue exclusion (unpublished data). Cells were then washed with TES/NaCl/EDTA buffer, pH 7.4, and transferred to Falcon 24 well plates (Becton Dickinson, Lincoln Park, NJ). Fluorescence was monitored in a Cytofluor II multiwell fluorescence plate reader (Perseptive Biosystems, Framingham, MA) with a quartz halogen lamp using 450 nm excitation/530nm emission or 560 nm excitation/620 nm emission wavelengths for *N*-NBD-PE or *N*-Rho-PE fluorescence, respectively. Liposome binding was determined as the amount of *N*-Rho-PE fluorescence associated with washed cells relative to total fluorescence of liposomes added. This percentage was converted to number of liposomes bound by multiplying by the number of liposomes added (assuming all liposomes were 100 nm in diameter and 10^5 lipid molecules/0.1 μ m diameter liposome. Therefore 6.02×10^{10} liposomes of 0.1 μ m diameter were added per sample). The process of lipid mixing is measured by the fluorescence dequenching (FDQ) of the NBD fluorophore. The % fluorescence dequenching (FDQ) was calculated by the following formula:

$$\left[\left(\frac{F_t}{F_{\max \text{ cells}}} \right) - \left(\frac{F_o \text{ alone}}{F_{\max \text{ alone}}} \right) \right] / \left[1 - \left(\frac{F_o \text{ alone}}{F_{\max \text{ alone}}} \right) \right] \times 100$$

where F_t = *N*-NBD-PE fluorescence of liposomes incubated with cells at a given time, F_o alone = initial *N*-NBD-PE fluorescence of liposomes only, $F_{\max \text{ cells}}$ and $F_{\max \text{ alone}}$ = maximal *N*-NBD-PE fluorescence of liposomes incubated either with cells or alone, respectively, as determined by addition of 0.5% C12E8

FDQ was assumed to result from all-or-none lipid mixing of liposomes with cells. Therefore % FDQ could be converted to number of liposomes mixed by simple multiplication of the total. This was done to take into account both the enhancement of binding and the lipid mixing after elastase activation.

Example 15**Optimum DODAP/MeO-suc-AAPV-DOPE composition for binding and lipid mixing with HL60 cells**

5 MeO-suc-AAPV-DOPE containing liposomes were designed to deliver their contents after binding, endocytic internalization, and fusion with and/or disruption of the endosomal membrane. DODAP was chosen instead of DOTAP (Pak et al., 1998) because only 20% of the DODAP population is positively charged at pH 7.4 (Bailey and Cullis, 1994). This allows a more complete peptide hydrolysis by elastase so that we can
10 model the more general case of cleavage on the liposome surface that is not affected by the high charge of the enzyme. For other enzymatic activators, it may be possible to utilize a permanently positively charged lipid that would allow the liposome to become positively charged at physiological pH after peptide cleavage such that extracellular binding would be triggered. By contrast, the tertiary amine of DODAP would be fully
15 protonated at pH 5, suggesting liposomes containing DODAP would undergo stronger interaction with negatively charged cell membranes within the low pH environment of the endocytic compartment.

To determine the optimum combination of DODAP and MeO-suc-AAPV-DOPE for triggerable binding and lipid mixing with cells the two lipids were formulated into
20 liposomes at different ratios. DODAP/MeO-suc-AAPV-DOPE liposomes prepared at 1:3, 1:1, and 3:1 mol ratios were pretreated with or without elastase and incubated with HL60 cells under low pH conditions to promote DODAP mediated binding to cells. Only DODAP/MeO-suc-AAPV₂-DOPE liposomes at a 1:1 mol ratio exhibited an elastase dependent increase in binding and lipid mixing with HL60 cells (Fig. 15), possibly as a
25 result of increased positive charge after enzymatic cleavage. The amount of DODAP in 1:3 liposomes was insufficient to mediate binding to cells, even at pH 5 after elastase treatment. By contrast, DODAP/MeO-suc-AAPV-DOPE (3:1 mol ratio) liposomes were able to bind to cells with or without elastase treatment, reflecting the greater amount of DODAP in these liposomes. These liposomes were also able to lipid mix with cells
30 without elastase activation. The DODAP/MeO-suc-AAPV-DOPE 1:1 mol ratio liposomal formulation was chosen for all further studies in order to develop a delivery system that can be triggered by enzymatic cleavage.

Example 16**Elastase-activated binding and lipid mixing of DODAP/MeO-suc-AAPV-DOPE liposomes with HL60 cells.**

5 DODAP was included in liposomes with MeO-suc-AAPV-DOPE to enhance binding with cells under low pH conditions. To determine whether the pH dependence of DODAP mediated binding is within physiological levels, DODAP/MeO-suc-AAPV-DOPE (1:1) liposomes were pretreated with elastase and incubated with HL60 cells at different pH. Enhanced binding and lipid mixing of elastase pretreated liposomes with
10 HL60 cells were observed when incubated at pH 4.6 or pH 5.1 (Fig. 16). Incubation at pH 5.8-7.4 did not yield any significant association of liposomes with cells. These results suggest that these liposomes are sensitive to elastase-mediated activation of binding and lipid mixing when DODAP is maximally positively charged. The pH required to achieve this state is present under normal physiological conditions with the late
15 endosome (Kielian et al., 1986).

To confirm that the enzymatic activity of elastase was responsible for triggering of binding and lipid mixing, DODAP/MeO-suc-AAPV-DOPE liposomes were pretreated with heat inactivated elastase. Heating elastase to 95 degrees C for 1 hour completely abrogated enzymatic cleavage of either a chromogenic substrate or MeO-suc-AAPV-
20 DOPE. Pretreatment of DODAP/MeO-suc-AAPV-DOPE liposomes with heat inactivated elastase did not enhance binding or lipid mixing with HL60 cells above background levels (Fig. 17). Only active elastase was capable of triggering this increased association. To observe this effect, pH 5 conditions were required, since at pH 7.4 the binding of these liposomes to HL60 cells was greatly reduced (Figs. 17A and 17B).

25

Example 17**Fluorescence microscopy of liposome-cell lipid mixing with HL60 cells**

DODAP/MeO-suc-AAPV-DOPE (1:1 mol/mol) liposomes were incubated for 2 hours at 37°C without or with elastase (5 • g protein/100 nmol lipid). Liposomes containing the
30 fluorescent lipid probes *N*-NBD-PE and *N*-Rho-PE were bound to HL60 cells in solution as described above. Cells were washed with HBSS to remove unbound liposomes and observed with a Bio-Rad (Hercules, CA) MRC-1000 laser scanning confocal imaging

system equipped with an Olympus BX50 microscope (Olympus Corp., Lake Success, NY) using an apochromat 60x oil (N.A. 1.40) objective. *N*-Rho-PE fluorescence was observed with the 568 nm line and observed at 605 nm emission. Images were frame averaged and false color was applied. All images within a figure were obtained under
5 identical conditions of confocal iris width, gain, and black level. Identical false color look up tables were also applied to images within a figure. Average fluorescence/cell μm^2 of all cells in an image was determined with the histogram feature of the Bio-Rad CoMOS confocal imaging software.

10 Confocal microscopy of liposomes mixed with HL60 cells confirmed that elastase pretreatment was required for enhanced binding and lipid mixing. Brightly fluorescently labeled cells indicate the *N*-Rho-PE fluorescent probe from the bound elastase-activated liposomes had mixed into the cell plasma membrane (Fig. 18). In contrast, untreated liposomes displayed significantly less binding and lipid mixing (Fig. 18). Quantitation of
15 fluorescence images revealed nearly 12 times as much fluorescence/cell area in HL60 cells that had been incubated with DODAP/MeO-suc-AAPV-DOPE liposomes treated with elastase, as compared to those cells incubated with untreated liposomes, although the distribution of this enhanced delivery may not be uniform across all the cells. Almost no fluorescence was associated with the cells at pH 7.4.

20

Example 18

Preparation of fluorescent dextran-containing liposomes and calcein-loaded liposomes

25 Fluorescent dextran containing liposomes were prepared by hydrating the DODAP/MeO-suc-AAPV-DOPE lipid film described previously (see Example 5) with a 50 mg/ml solution of 10,000 MW tetramethyl rhodamine dextran (TMR-dextran) in TES/NaCl/EDTA buffer. Liposomes were then vortexed, freeze/thawed, and extruded through 0.1 μm filters as described above. To remove unencapsulated dextran the
30 liposome solution was extensively dialyzed with TES/NaCl/EDTA buffer using a Biodialyser (Sialomed, Columbia, MD) fitted with 50 nm pore size filters. Calcein-loaded liposomes

were prepared by hydrating the lipid film in the presence of buffer containing 50mM calcein. Calcein (> 95% pure) and tetramethylrhodamine dextran were obtained from Molecular Probes (Eugene, OR). Calcein solution was adjusted to a pH of approximately 7 and an osmolarity of approximately 300 mosm prior to preparation of
5 liposomes. Calcein-loaded liposomes contained 0.75 mol % N-Rho-PE to monitor liposome binding. After preparation of large vesicles as described above, calcein-loaded liposomes were transferred to a 10,000 molecular weight cut off (MWCO) Slide-A-Lyzer (Pierce, Rockford, IL) and extensively dialyzed with TES/NaCl/EDTA buffer. The encapsulated volume of these liposomes was 0.8 l/mol of lipid. Sonicated vesicles
10 were prepared by drying lipid in the same manner as described above but preparations were vortexed then water bath sonicated for >10 min at room temperature. Lipid concentration was monitored by phosphate assay (Bartlett 1959). The size of liposomes was verified by quasi-elastic light scattering using a Nicomp Submicron Particle Sizer (Particle Sizing Systems, Santa Barbara, CA). Freeze-thaw/extrusion
15 vesicles and sonicated vesicles were 70-80 nm and 35-45 nm in diameter, respectively, as determined by number weighted Gaussian analysis.

Example 19

Aqueous contents delivery from DODAP/MeO-suc-AAPV-DOPE liposomes to HL60 20 cells

To determine if the enhanced lipid mixing between elastase-activated DODAP/MeO-suc-AAPV-DOPE liposomes and HL60 cells is truly indicative of fusion, the delivery of an aqueous probe from the liposome to the cell
cytoplasm was monitored. DODAP/MeO-suc-AAPV-DOPE liposomes were loaded with
25 tetramethyl rhodamine labeled 10,000 MW dextran (TMR-dextran), treated with or without elastase, and incubated with HL60 cells under pH 5 conditions.

TMR-dextran loaded DODAP/MeO-suc-AAPV-DOPE liposomes (40 nmol, prepared as described in Examples 5 and 16) were incubated with 1×10^5 HL60 cells in
30 $200 \mu\text{l}$ TES/NaCl/EDTA buffer under pH 5, 37°C, conditions for 30 min to induce binding. TMR-dextran fluorescence was observed by confocal microscopy under the same conditions as N-Rho-PE described in Example 17.

Only DODAP/MeO-suc-AAPV-DOPE liposomes that had been pretreated with elastase were capable of fusing with HL60 cells, as demonstrated by TMR-dextran labeling of the cytoplasm of these cells (Fig. 19). HL60 cells incubated with liposomes that had not been treated with elastase contained little or no cytoplasmic fluorescent dextran, indicating elastase cleavage was required to trigger the fusion of DODAP/MeO-suc-AAPV-DOPE liposomes with HL60 cells.

TMR-dextran delivery to cells was not due to leakage of the fluorescent dextran out of the liposomes and subsequent uptake by HL60 cells. This possibility was investigated by inducing TMR-dextran release from DODAP/MeO-suc-AAPV-DOPE liposomes. The results demonstrated that there was no uptake into the HL60 cells.

Example 20

15 DODAP/MeO-suc-AAPV-DOPE liposome interaction with EVC304 cells: binding, lipid mixing and calcein delivery

Liposomes were bound to adherent ECV304 cells via a biotin-streptavidin linkage. To this end DODAP/MeO-suc-AAPV-DOPE (1:1 mol:mol) liposomes were prepared with 0.3 mol% N-biotinyl caproylamine-PE (N-biotinyl-cap-PE) as well as fluorescent lipid probes or with encapsulated calcein. ECV304 cells that had been plated on glass coverslips in tissue culture plates were washed with HBSS buffer and then incubated sequentially at room temperature with biotin-wheat germ agglutinin (WGA) (20 • g/ml, obtained from Pierce, Rockford, IL) and streptavidin (40 • g/ml, obtained from Molecular Probes, Eugene, OR) prepared in HBSS, 30 min/treatment. Cells were washed after each treatment. Liposomes were treated with or without elastase as described above. Certain aliquots of pretreated DODAP/MeO-suc-AAPV-DOPE (1:1 mol/mol) liposomes were freeze/thawed after dialysis and prior to addition to cells to release the liposomal contents. Such freeze/thawed liposomes were exposed to liquid nitrogen/37°C water bath for 5 cycles. Self-quenching of calcein was reduced by approximately 85% (maximal FDQ determined by detergent solubilization) after freeze/thawing, indicating release of encapsulated calcein. In all cases, 50-100 nmol of liposomes were added to confluent ECV304 cell monolayers (1x10⁵ cells/well of 24 well

plate) and incubated in HBSS for 30 min at room temperature to promote *N*-biotinyl cap-PE binding to streptavidin. Unbound liposomes were removed by repeated washes. After the final wash, fresh HBSS buffer was added to all wells and cells were incubated at 37°C for given times. Fluorescence was quantitated as described above. Calcein
5 fluorescence was excited with the 488 nm line of a krypton/argon laser and observed at 522 nm emission

Example 21

Elastase-activated binding and, lipid mixing of DODAP/MeO-suc-AAPV-DOPE 10 liposomes with adherent ECV304 cells

To determine whether elastase-treated liposomes internalized within mammalian cells by endocytosis could be activated to bind and fuse with the cells, liposomes were bound to cells under physiological conditions at pH 7.4 via a biotin-streptavidin linkage. The pretreatment of DODAP/MeO-suc-AAPV-DOPE liposomes with elastase activated
15 these liposomes to bind and fuse with HL60 cells when the pH was artificially lowered to mimic the endosomal environment (Fig. 19). The adherent cell line, ECV304 was selected to circumvent complications arising from biotin-streptavidin mediated aggregation of the suspension HL60 cells.

ECV304 cells were sequentially treated with biotinylated-wheat germ agglutinin
20 and streptavidin. DODAP/MeO-suc-AAPV-DOPE liposomes containing trace amounts of *N*-biotinyl cap-PE and pretreated with or without elastase were then added to the cells and incubated at pH 7.4. Elastase pretreated liposomes exhibited enhanced binding, as compared to untreated liposomes (Fig. 20), perhaps as a result of decreased negative surface charge or better accessibility of the biotinyl group. It appears that elastase
25 activation augments the biotin-streptavidin mediated binding. The biotin-streptavidin linkage was required, as the absence of biotin-streptavidin treatment abrogated any binding of untreated or elastase pretreated liposomes at this pH (unpublished data). Liposomes with or without elastase activation were localized within perinuclear endocytic vesicles (Fig. 21). There was enhanced binding of elastase pretreated liposomes and
30 uptake into endocytic vesicles; however, there was only a slight increase in lipid mixing of these liposomes with ECV304 cells (Fig. 20). This may be due to the nature of the assay because the fluorescence dequenching assay requires the diffusion of the *N*-

NBD-PE and *N*-Rho-PE probe after lipid mixing. Diffusion of the fluorescent lipid probes after fusion with the endosomal membrane may be insufficient to completely diminish the resonance energy transfer. Therefore an aqueous contents delivery assay was employed.

5

Example 22

Aqueous contents delivery of calcein from DODAP/MeO-suc-AAPV-DOPE liposomes to adherent ECV304 cells

DODAP/MeO-suc-AAPV-DOPE liposomes were loaded with self-quenched
10 concentrations of calcein, a fluorescent aqueous probe, as previously described (Examples 5 and 18). After extensive dialysis to remove unencapsulated calcein these liposomes were treated with or without elastase and bound to ECV304 cells by biotinylated WGA-streptavidin. Elastase pretreated liposomes displayed relatively rapid calcein dequenching that increased over the course of several hours (Fig. 22). Maximal
15 dequenching of calcein appears to be achieved after 2 hours at 37°C, which was consistent with the time course of DODAP/MeO-suc-AAPV-DOPE liposome endocytosis (unpublished data) and cationic lipid:DNA complex uptake [Zabner et al., 1995]. The apparent delivery of calcein was completely dependent upon liposome-cell interactions, as calcein-loaded liposomes subjected to identical conditions of elastase pretreatment
20 and pH 5 environment but without co-incubation with cells did not demonstrate any increase in calcein fluorescence dequenching. DODAP/MeO-suc-AAPV-DOPE liposomes that had not been treated with elastase exhibited much less initial calcein dequenching that slowly increased over time, albeit not to levels observed with the elastase-activated liposomes (Fig. 22). Repeated freeze/thaw cycles of DODAP/MeO-
25 suc-AAPV-DOPE liposomes results in almost complete release of encapsulated calcein. Incubation of this mixture of freeze/thawed liposomes and released free calcein with ECV304 cells resulted in only 2-3% of the calcein uptake observed with cells incubated with intact liposomes. Thus the fluorescence dequenching observed with elastase treated intact liposomes was due to delivery of calcein to the cells rather than a non-
30 specific uptake of free calcein. The results with the adherent endothelial cell line ECV304 are similar to those obtained with the HL60 leukemia cell line described in earlier examples.

Confocal microscopy of ECV304 cells incubated with elastase pretreated DODAP/MeO-suc-AAPV-DOPE liposomes contained brightly fluorescent perinuclear localized vesicles that appear to be due to uptake of the calcein-loaded liposomes into
5 endosomes. Importantly, these cells also had diffuse fluorescence within the cells (Fig. 21) that is indicative of calcein delivery into the cytosol. The diffuse fluorescence was not due to non-specific uptake of calcein released from liposomes prior to endocytosis, as loaded liposomes subjected to freeze/thaw cycles did not result in fluorescently labeled cells. Unactivated liposomes were also endocytosed into vesicles near the
10 nucleus. However, there was no diffuse fluorescence visible with these liposomes. Quantitation of total fluorescence/cell area showed ECV304 cells incubated with elastase-activated DODAP/MeO-suc-AAPV-DOPE liposomes had more than twice the amount of calcein fluorescence as those incubated with unactivated liposomes. Without
15 - elastase pretreatment there does not appear to be significant delivery of the encapsulated calcein into the cell cytoplasm.

Example 23**Abbreviations**

	ATCC	American Type Culture Collection
5	BSA	Bovine serum albumin
	DCC	Dicyclohexyl carbodiimide
	DC-Chol	3-beta-[N-[(N'.N'-dimethylamino)ethane]carbamoyl]cholesterol
	DCU	Dicyclohexyl urea
	DMRI	1,2-dimyristooxypropyl-3-dimethylhydroxyethyl ammonium bromide
10	DODAP	1-N,N-dimethylamino dioleoyl propane
	DOPE	Dioleoyl phosphatidylethanolamine
	DORIE	1,2-dioleooxypropyl-3-dimethylhydroxyethyl ammonium bromide
	DOTAP	1,2-bis(oleoyloxy)-3-trimethylammonio)propane
	DPPE	Dipalmityl phosphatidylethanolamine
15	DSPE	Distearoyl phosphatidylethanolamine
	EDTA	Ethylenediamine tetraacetic acid
	EGTA	Ethylenebis(oxyethylenenitrilo)-tetraacetic acid
	FBS	Fetal bovine serum
	FDQ	Fluorescence dequenching
20	HBSS	Hanks' Balanced Salts solution
	HLE	Human leukocyte elastase
	LUV	Large Unilamellar vesicles
	MeO-suc	Methoxy succinyl
	MLV	Multilamellar vesicles
25	N-biotinyl-cap-PE	N-biotinyl caproylamine-PE
	PC	Phosphatidylcholine
	PE	Phosphatidylethanolamine
	POPE	Palmitoyl oleoyl phosphatidylethanolamine
	PS	Phosphatidylserine
30	RBC	Red blood cell
	SUV	Small unilamellar vesicles
	TFA	Trifluoroacetic acid

THF	Tetrahydrofuran
TLC	Thin layer chromatography
TMR-dextran	Tetramethyl rhodamine dextran
tPE	Transesterified phosphatidylethanolamine
5 WGA	Wheat germ agglutinin.

References:

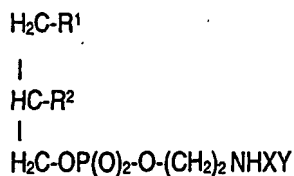
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What is claimed is:

1. A method of administering the contents of a liposome to a mammal which comprises administering to the mammal a composition comprising:
 - (i) a pharmaceutically acceptable carrier; and
 - (ii) a liposome comprising a lipid component which comprises a peptide-lipid conjugate having the formula:



wherein:

X is a linker selected from the group consisting of a single bond or the group R³; each of R¹ and R² is

$-\text{OC}(\text{O})(\text{CH}_2)_{n1}(\text{CH}=\text{CH})_{n2}(\text{CH}_2)_{n3}(\text{CH}=\text{CH})_{n4}(\text{CH}_2)_{n5}(\text{CH}=\text{CH})_{n6}(\text{CH}_2)_{n7}(\text{CH}=\text{CH})_{n8}(\text{CH}_2)_{n9}\text{CH}_3$; and

R³ is

$-\text{C}(\text{O})(\text{CH}_2)_{n1}(\text{CH}=\text{CH})_{n2}(\text{CH}_2)_{n3}(\text{CH}=\text{CH})_{n4}(\text{CH}_2)_{n5}(\text{CH}=\text{CH})_{n6}(\text{CH}_2)_{n7}(\text{CH}=\text{CH})_{n8}(\text{CH}_2)_{n9}\text{HN}-$.

n₁ is zero or is an integer equal to from 1 to 22,

n₃ is zero or is an integer equal to from 1 to 19,

n₅ is zero or is an integer equal to from 1 to 16,

n₇ is zero or is an integer equal to from 1 to 13, and

n₉ is zero or is an integer equal to from 1 to 10;

for each of R¹ and R² the sum of n₁ + 2n₂ + n₃ + 2n₄ + n₅ + 2n₆ + n₇ + 2n₈ + n₉ is an integer equal to from 12 to 22;

for R³ the sum of n₁ + 2n₂ + n₃ + 2n₄ + n₅ + 2n₆ + n₇ + 2n₈ + n₉ is zero or is an integer equal to from 1 to 22;

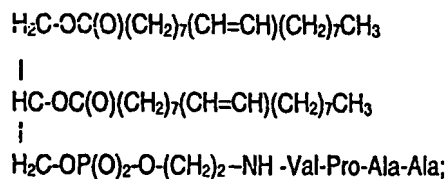
each of n₂, n₄, n₆ and n₈ is equal to 0 or 1;

Y is a peptide comprising an amino acid sequence which is the substrate of a cell-secreted or cell-associated peptidase; and

the contents of the liposome are delivered to the vicinity of cells in the mammal which secrete a peptidase which recognizes the amino acid substrate.

2. The method of claim 1, wherein the liposome is a large unilamellar liposome.
3. The method of claim 1, wherein the peptide comprises the amino acid sequence Ala-Ala.
4. The method of claim 3, wherein the peptide comprises an amino acid sequence selected from the group consisting of the sequences Ala-Ala, Ala-Ala-Pro-Val (SEQ ID NO: 1), Ala-Ala-Met-, Ala-Ala-Pro-Phe (SEQ ID NO: 3), Ala-Ala-Pro-Met (SEQ ID NO: 4), Ala-Ala-Arg, Ser-Ala-Ala-Arg (SEQ ID NO: 5), Ser-Ser-Ala-Ala-Arg (SEQ ID NO: 6), Ser-S carboxyl sugar-Ala-Ala-Arg- (SEQ ID NO: 7), Ala-Ala-Asp-, Ser-Ala-Ala-Asp- (SEQ ID NO: 8) and Ser-Ser-Ala-Ala-Asp- (SEQ ID NO: 9).
5. The method of claim 4, wherein the peptide comprises the amino acid sequence Ala-Ala-Pro-Val (SEQ ID NO: 1).
6. The method of claim 1, wherein the peptide comprises an amino acid sequence selected from the group consisting of Arg-Pro-Lys-Pro-Leu-Ala-Nva- (SEQ ID NO: 10), Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva- (SEQ ID NO: 11), Ser-Ser-Arg-Pro-Lys-Pro-Leu-Ala-Nva- (SEQ ID NO: 12), Pro-Cha-Gly-Nva-His-Ala-Dpa-NH₂ (SEQ ID NO: 13), Pro-Leu-Gly-Leu-Dpa-Ala-Arg-NH₂ (SEQ ID NO: 14), ; Pro-Cha-Gly-Nva-; Pro-Leu-Gly-Leu- (SEQ ID NO: 15) and Gly-Pro-Gln-Gly-Ile- (SEQ ID NO: 16).
7. The method of claim 1, wherein the peptide is modified at its amino terminus by a moiety selected from the group consisting of acetyl, methoxy, carboxy sugar, polyethylene glycol and methoxy-substituted carboxy sugar modifications.
8. The method of claim 7, wherein the peptide is preferably N-acetyl modified at its amino terminus and the peptide-lipid conjugate comprises from about 20 mole % to about 80 mole % of the lipid component.
9. The method of claim 1, wherein X is a single bond, each of R¹ and R² is -OC(O)(CH₂)₇(CH=CH)(CH₂)₇CH₃ and the peptide comprises the amino acid sequence Ala-Ala.

10. The method of claim 9, wherein the peptide-lipid conjugate is:



the peptide is N-methoxysuccinyl modified at its amino terminus and the peptide-lipid conjugate comprises from about 20 mole % to about 80 mole % of the lipid component.

11. The method of claim 10, wherein the lipid component further comprises an additional lipid which is a positively charged lipid selected from the group consisting of 1,2-bis(oleoyloxy)-3(trimethylammonio)propane (DOTAP); 1-N,N-dimethylamino dioleoyl propane (DODAP); 1-oleoyl-2-hydroxy-3-N,N-dimethylamino propane; 1,2-diacyl-3-N,N-dimethylamino propane; 1,2-didecanoyl -1-N,N,-dimethylamino propane, 3-beta-[N-[(N',N'-dimethylamino)ethane]carbamoyl]cholesterol (DC-Chol), 1,2-dimyristooxypropyl-3-dimethylhydroxyethyl ammonium bromide (DMRIE); and 1,2-dioleooxypropyl-3-dimethylhydroxyethyl ammonium bromide (DORI).
12. The method of claim 11, wherein the positively charged lipid is DODAP.
13. The method of claim 12, wherein the lipid component comprises 50 mole % DODAP and 50 mole % of the peptide-lipid conjugate.
14. The method of claim 12, wherein the lipid component comprises DODAP and N-Ala-Ala-Pro-Val-DOPE.
15. The method of claim 11, wherein the positively charged lipid is DOTAP.
16. The method of claim 15, wherein the lipid component comprises 50 mole % DOTAP and 50 mole % of the peptide-lipid conjugate.
17. The method of claim 15, wherein the lipid component comprises 50 mole % DOTAP and 50 mole % N-Ala-Ala-Pro-Val.

18. The method of claim 10, wherein the lipid component further comprises an additional lipid selected from the group consisting of trans-esterified phosphatidylethanolamine (tPE), dipalmitoyl phosphatidylethanolamine (DPPE), palmitoyl oleoyl phosphatidylethanolamine (POPE) and dioleoyl phosphatidylethanolamine (DOPE).
19. The method of claim 1, wherein the peptidase is selected from the group consisting of elastase, plasmin, plasminogen activator, urokinase; stromelysin, human collagenases, cathepsins, lysozyme, granzymes, dipeptidyl peptidases, peptide hormone-inactivating enzymes, kininases, bacterial peptidases and viral proteases.
20. The method of claim 19, wherein the peptidase is elastase.
21. The method of claim 19, wherein the peptidase is stromelysin, a cathepsin, plasmin or a plasminogen activator.
22. The method of claim 1, wherein the liposome further comprises a bioactive agent selected from the group consisting of antiviral agents, antibacterial agents, antifungal agents, antineoplastic agents, antiinflammatory agents, radiolabels, radiopaque compounds, fluorescent compounds, mydriatic compounds, bronchodilators, local anesthetics, nucleic acid sequences, plasmid deoxyribonucleic acid sequences and bioactive lipids.
23. The method of claim 22, wherein the bioactive agent is an anticancer agent, the liposome contains a therapeutically effective amount of the anticancer agent and the mammal is afflicted with a cancer.
24. The method of claim 23, wherein the cancer is selected from the group consisting of brain cancers, breast cancers, carcinomas, colon cancers, leukemias, lung cancers, lymphomas, ovarian cancers and sarcomas.
25. The method of claim 22, wherein the bioactive agent is an anti-inflammatory agent, the liposome comprises a therapeutically effective amount of the anti-inflammatory agent and the mammal is afflicted with an inflammatory disorder.
26. The method of claim 22, wherein the bioactive agent is a nucleic acid, the mammal is afflicted with a genetic disorder and the nucleic acid encodes a protein which ameliorates the disorder.

Fig. 1

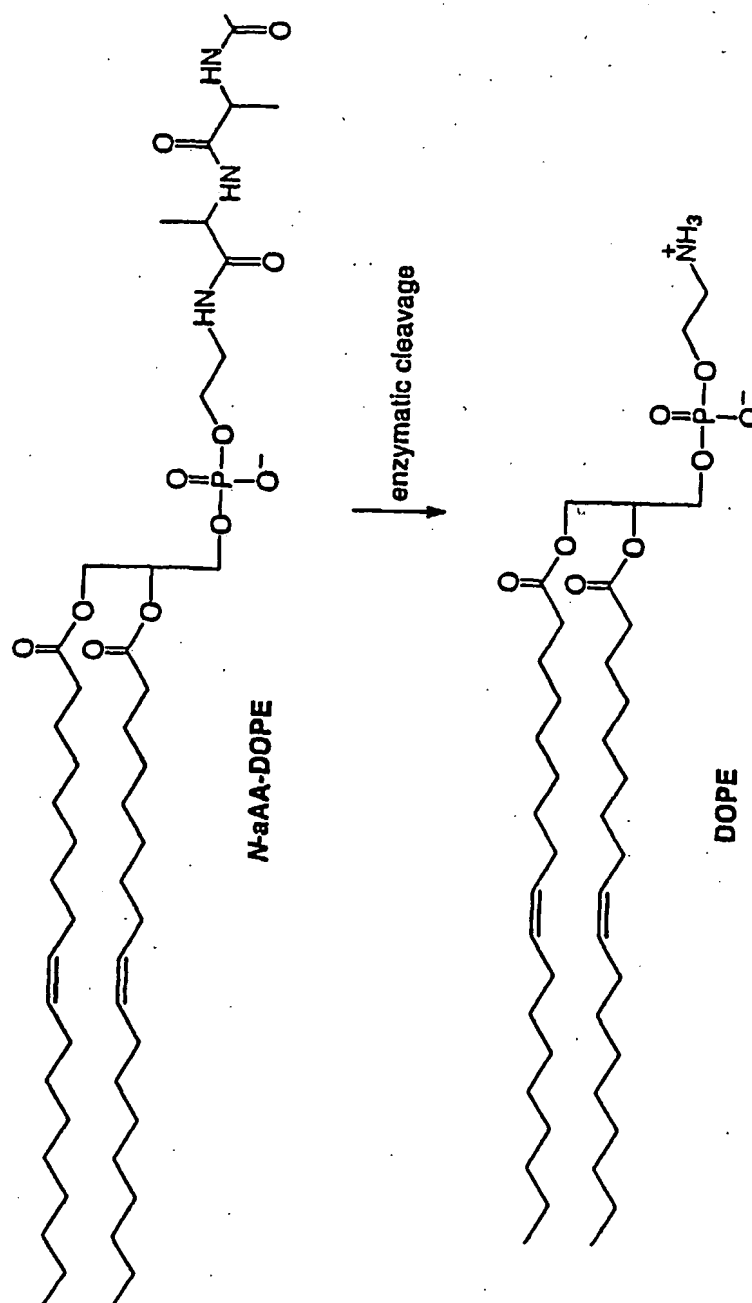
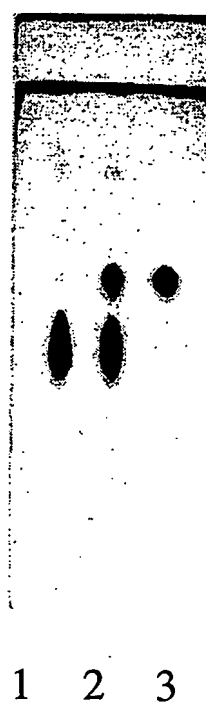


Fig. 2
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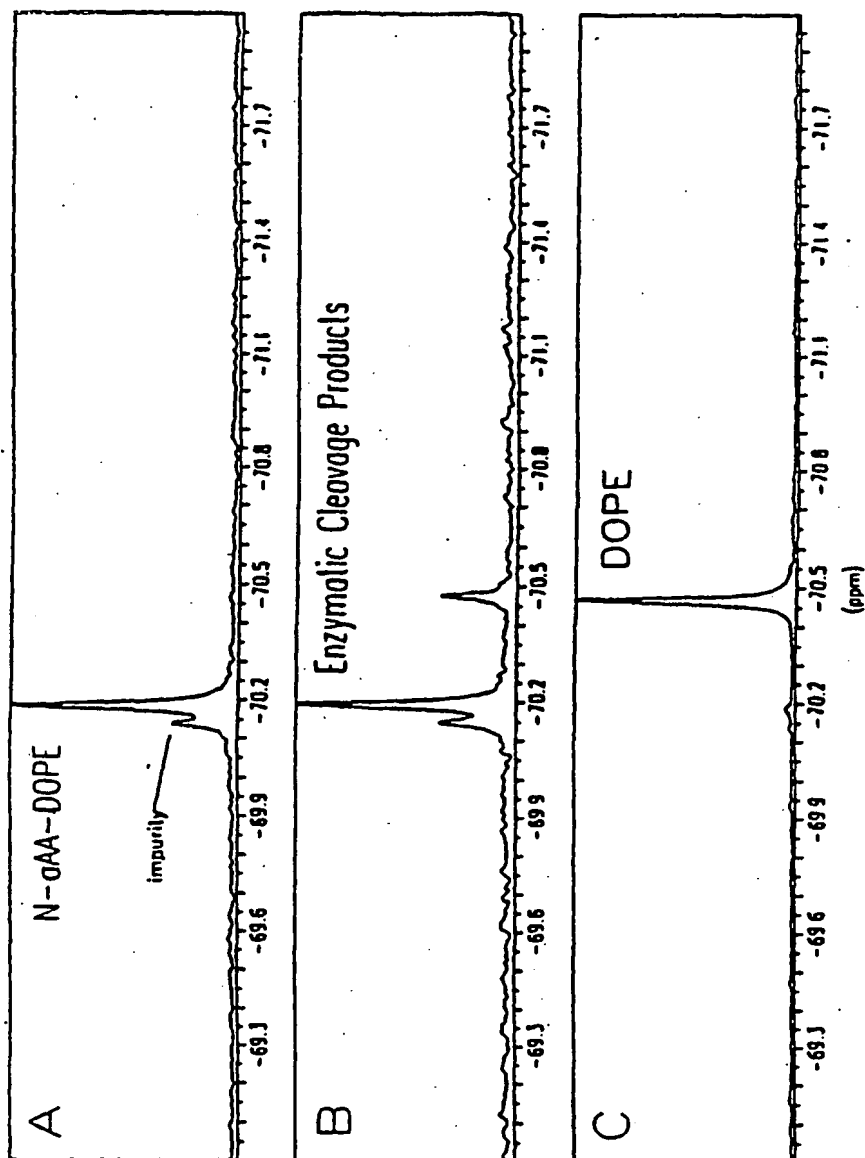


Fig. 2
B



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Fig. 3



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Fig. 4

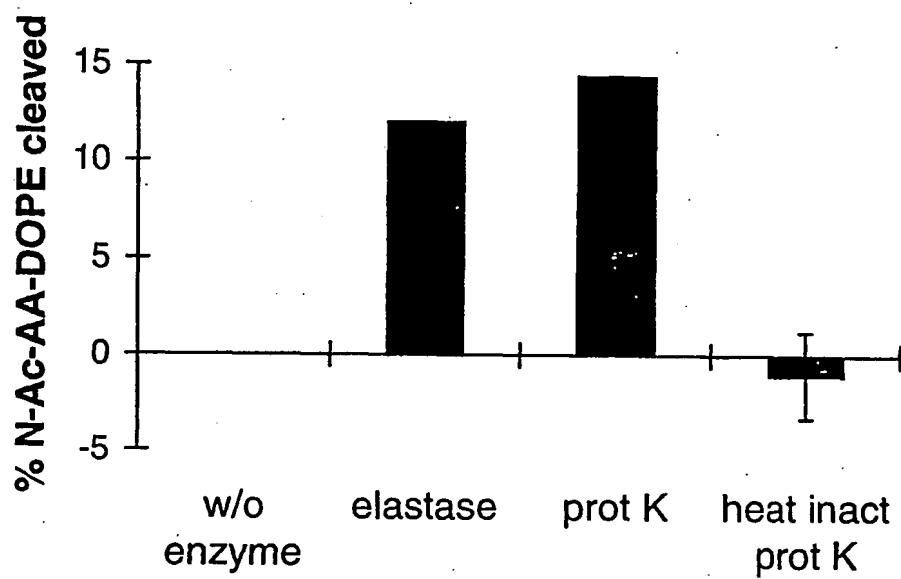
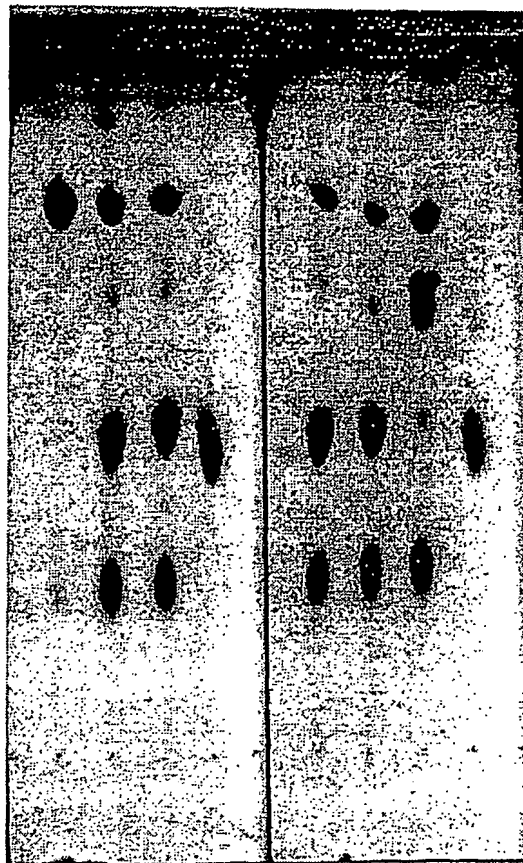


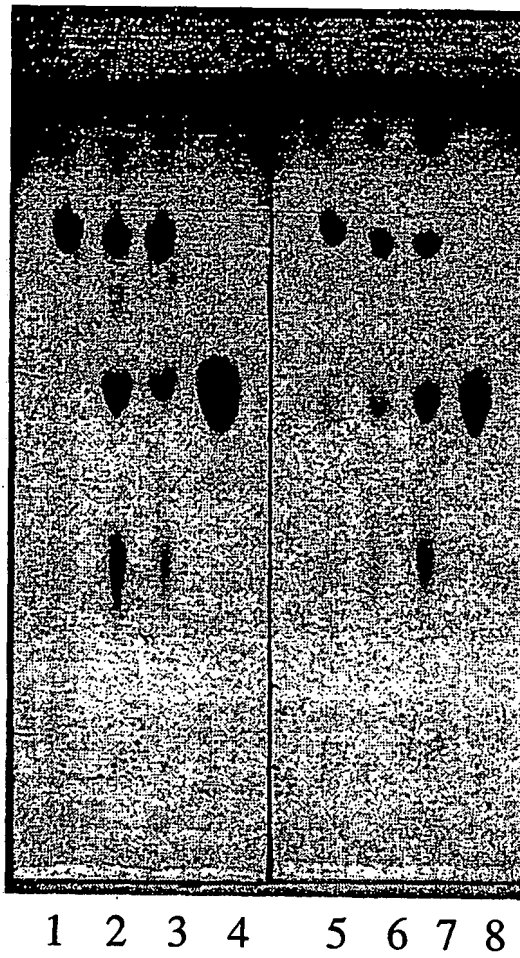
Fig. 5A



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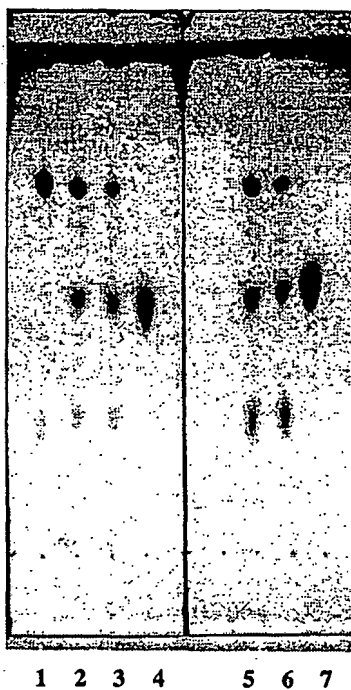
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Fig. 5B



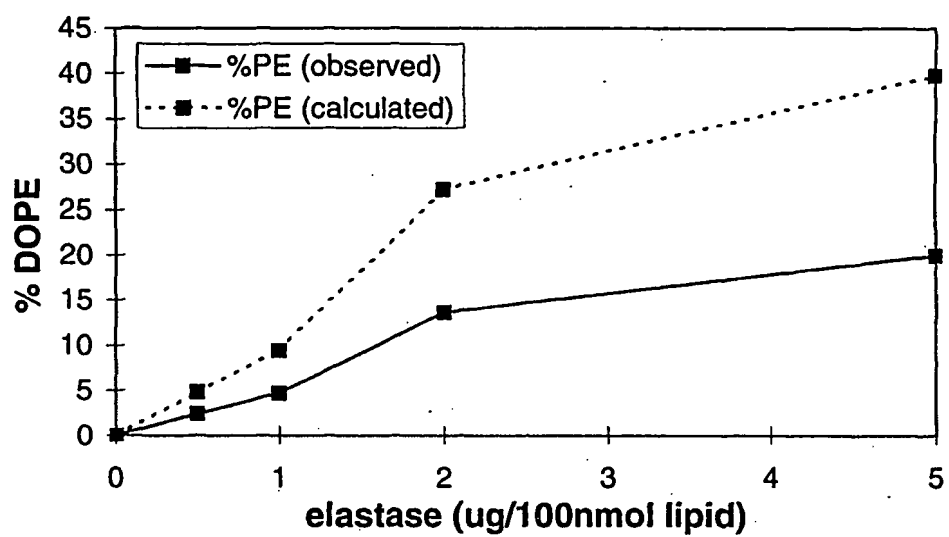
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Fig. 5C



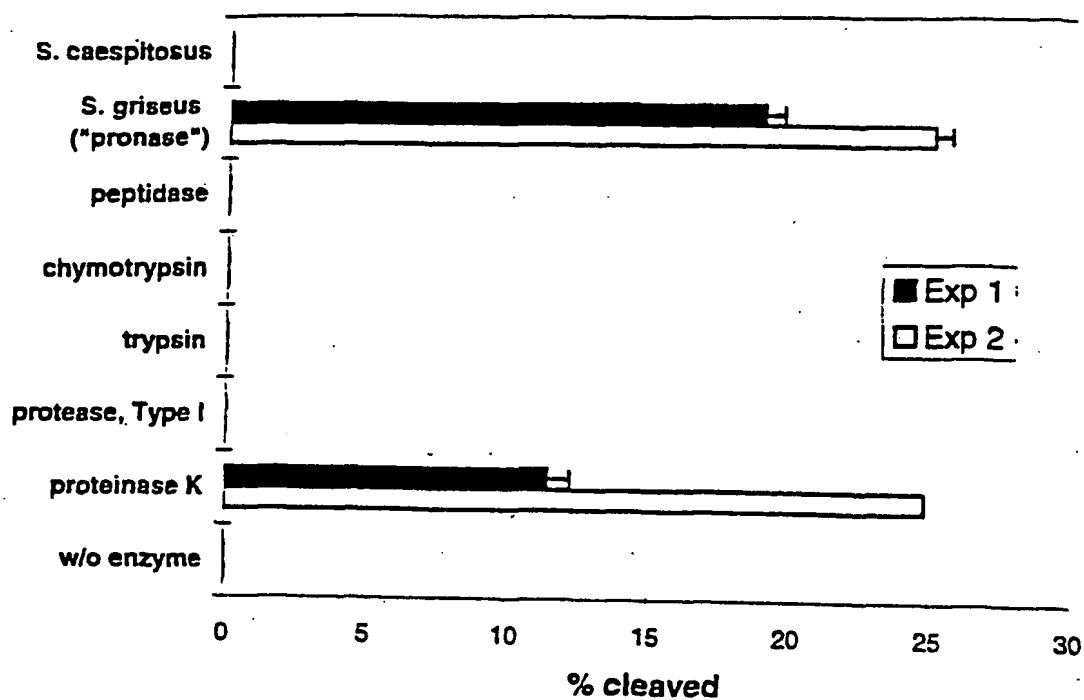
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Fig. 6



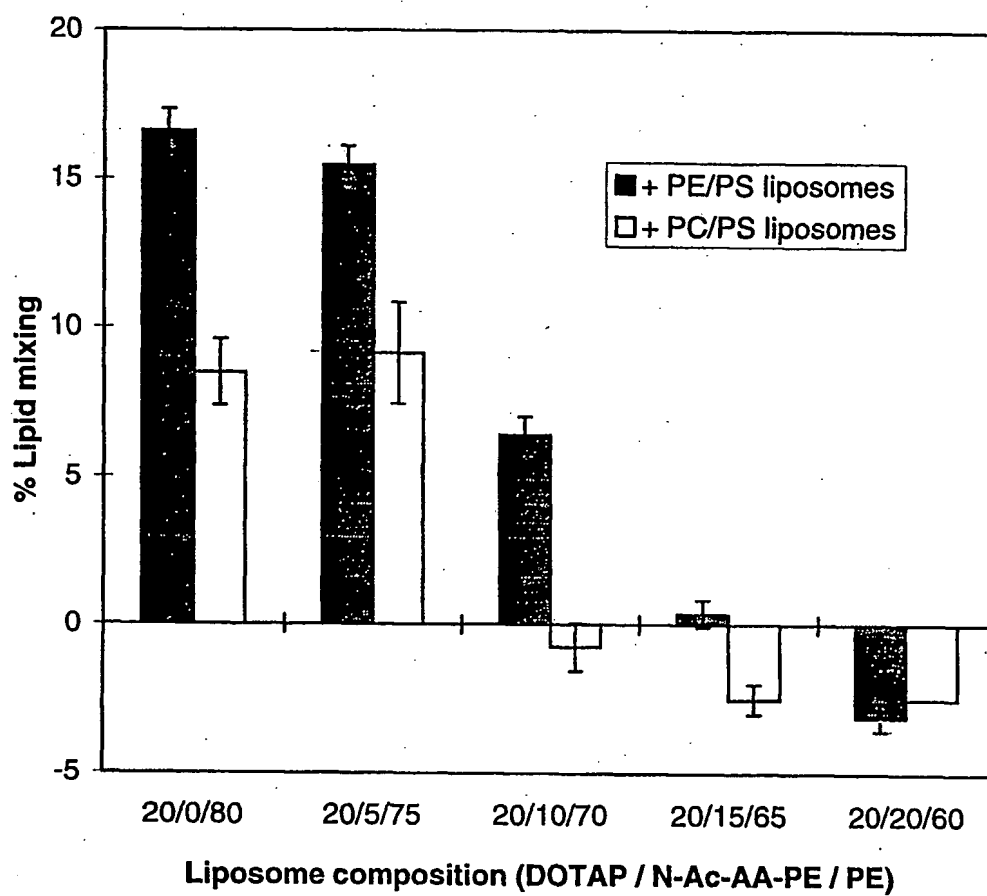
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Fig. 7



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Fig. 8A



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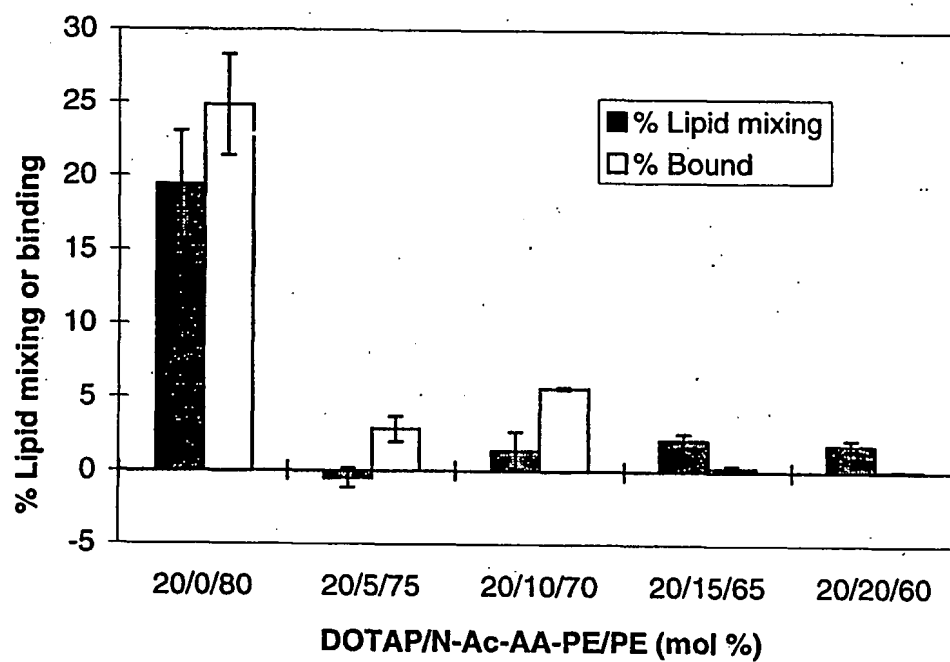
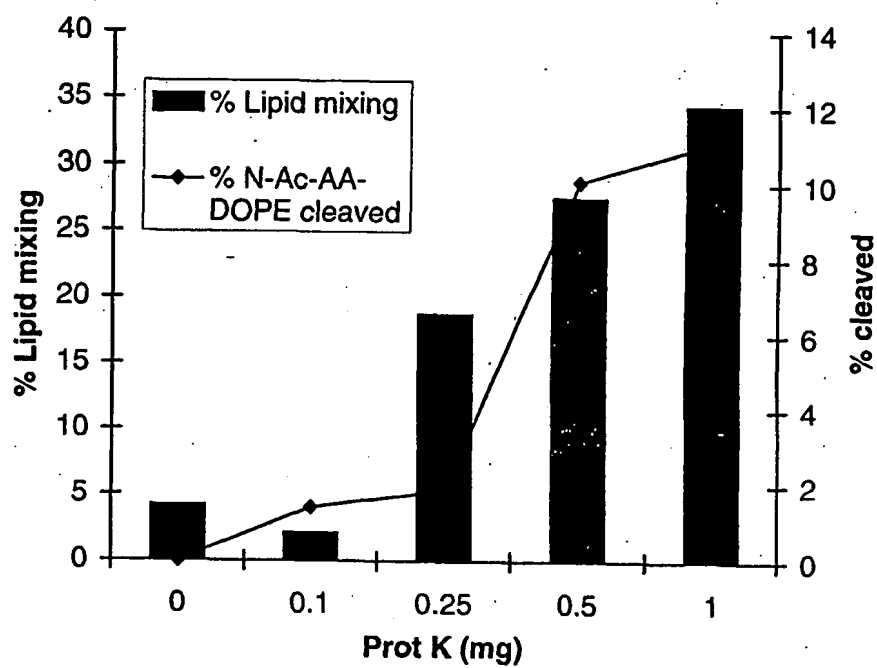


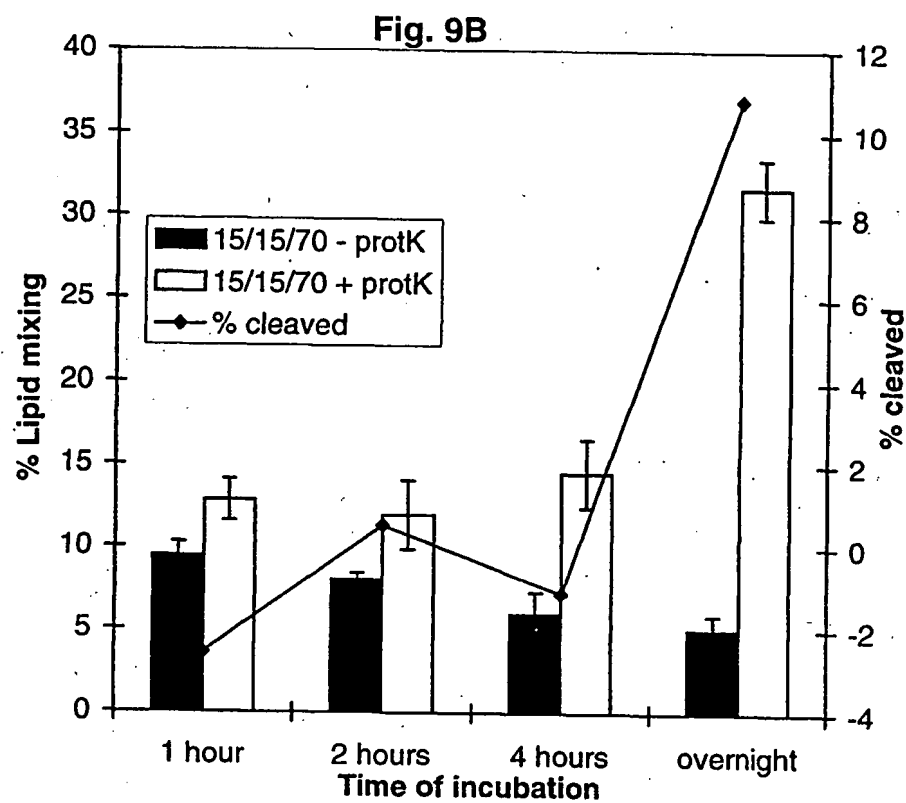
Fig. 8B

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Fig. 9A

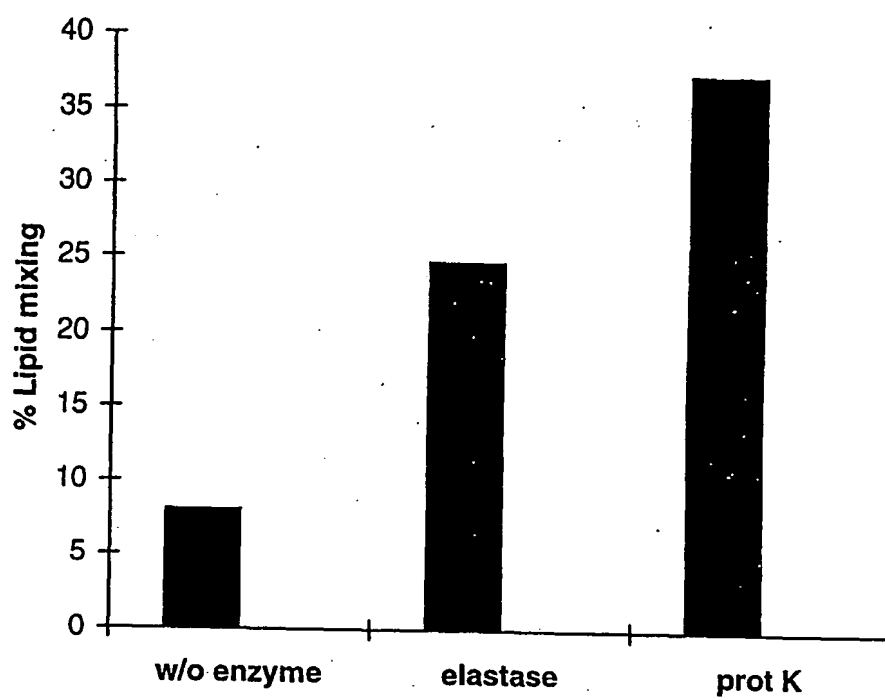


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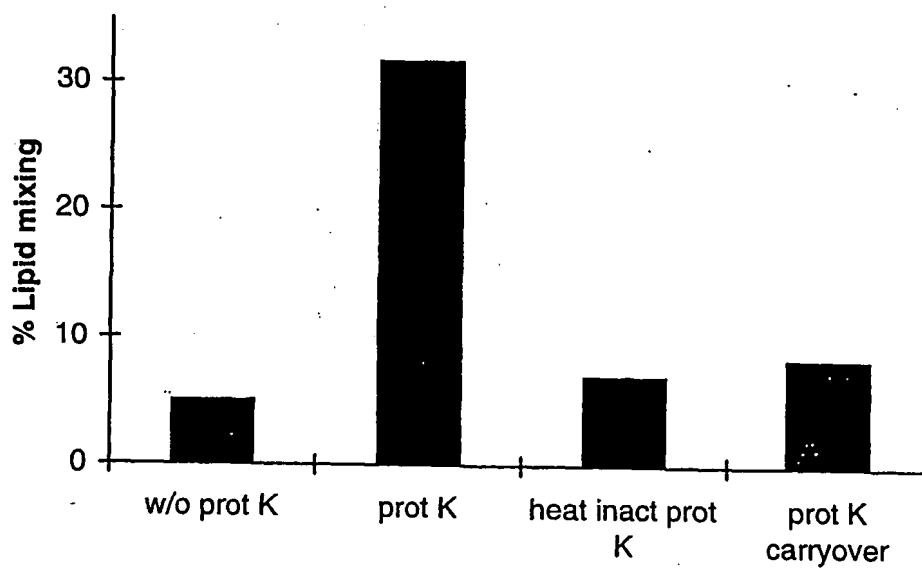
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Fig. 10



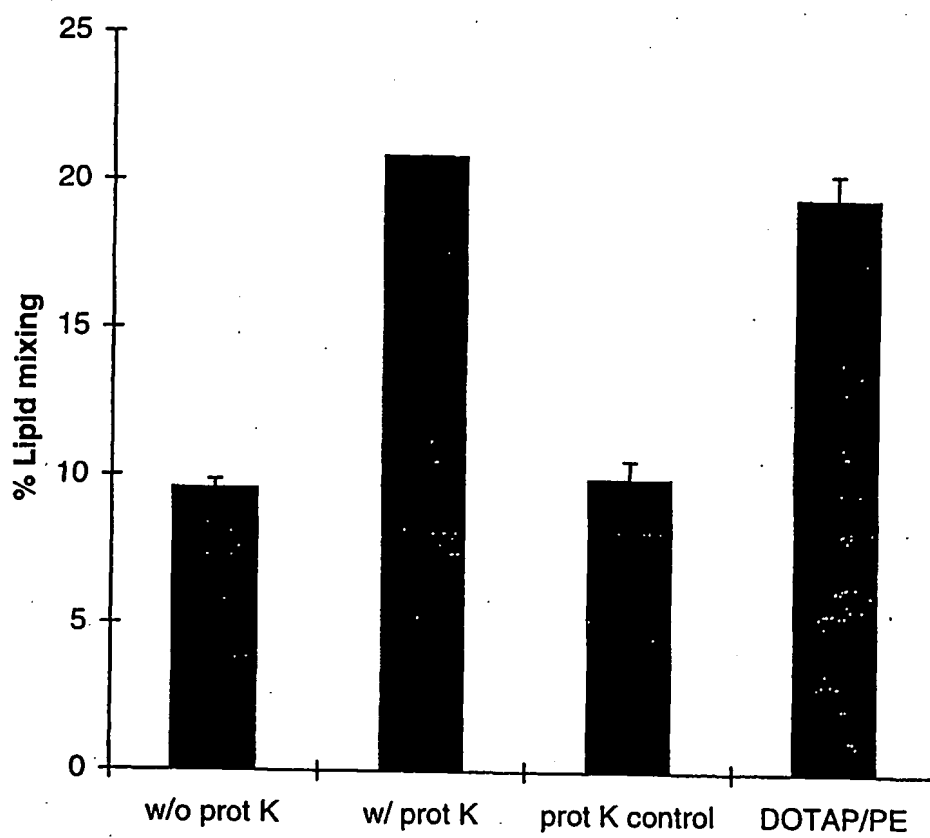
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Fig. 11



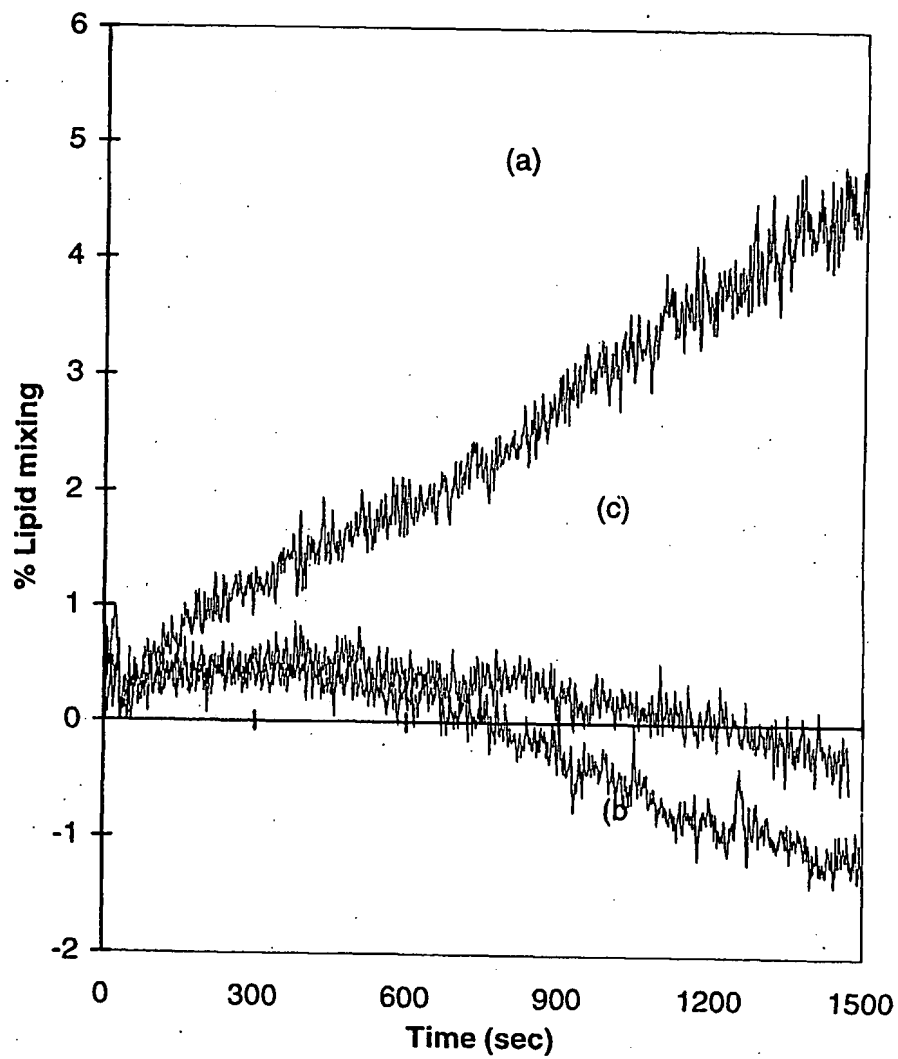
16/30

Fig. 12



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Fig. 13



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Fig. 14A

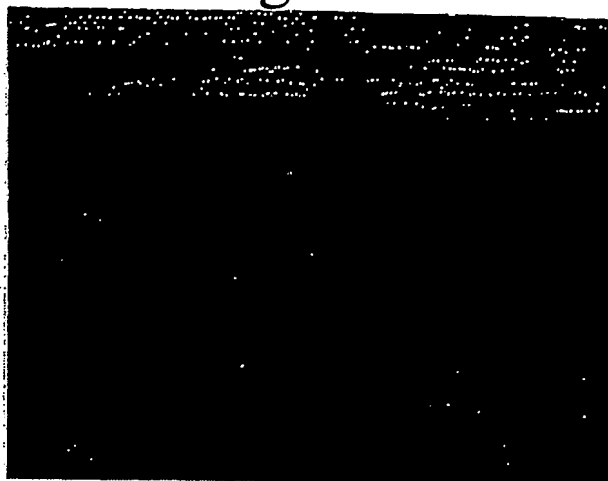


Fig. 14B



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Fig. 14C

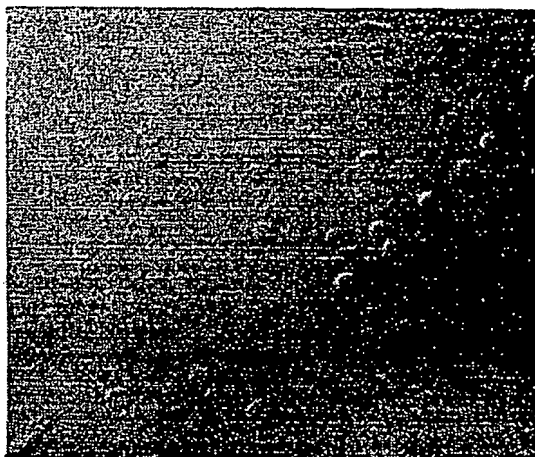
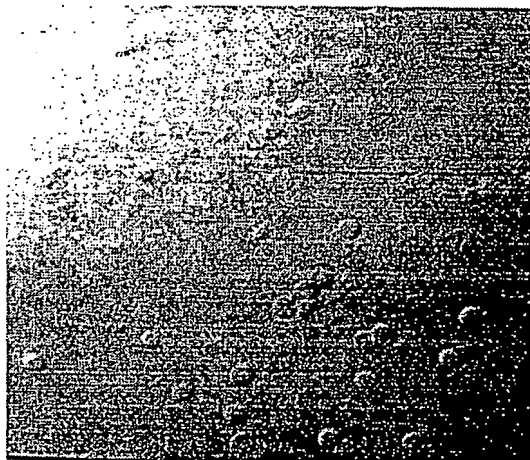


Fig. 14D



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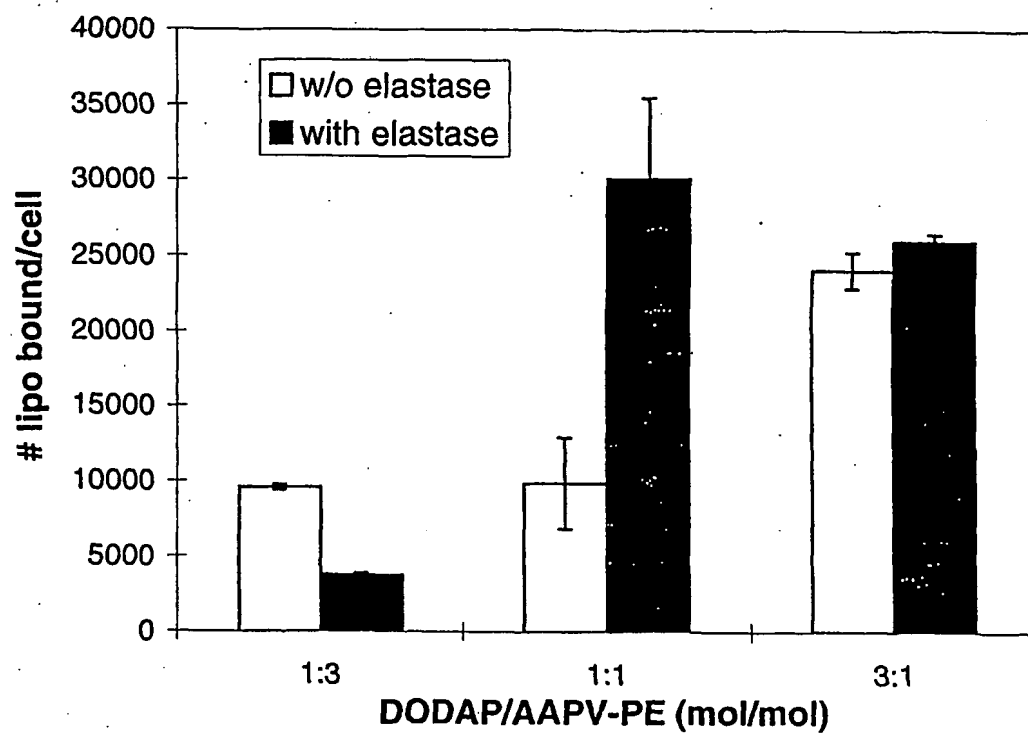


Fig. 15A

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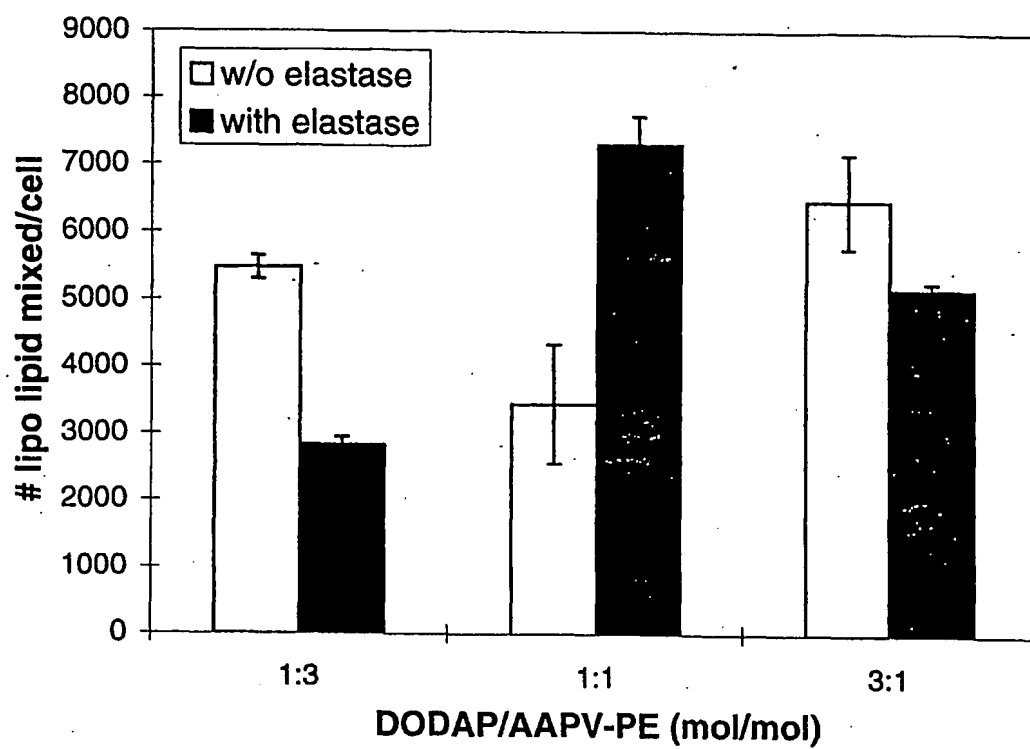


Fig. 15B

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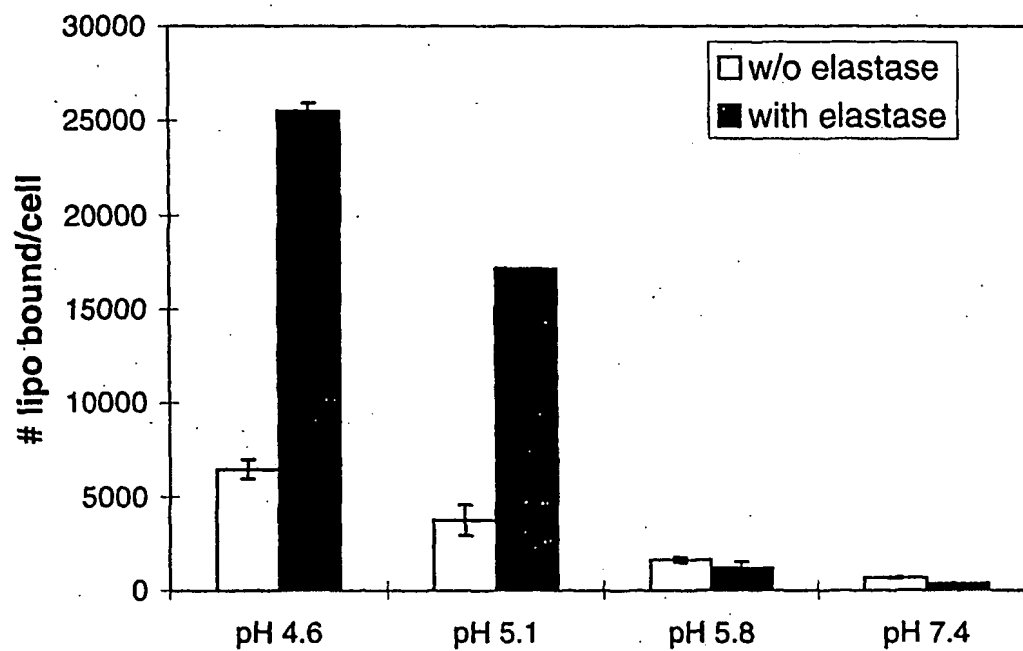


Fig. 16A

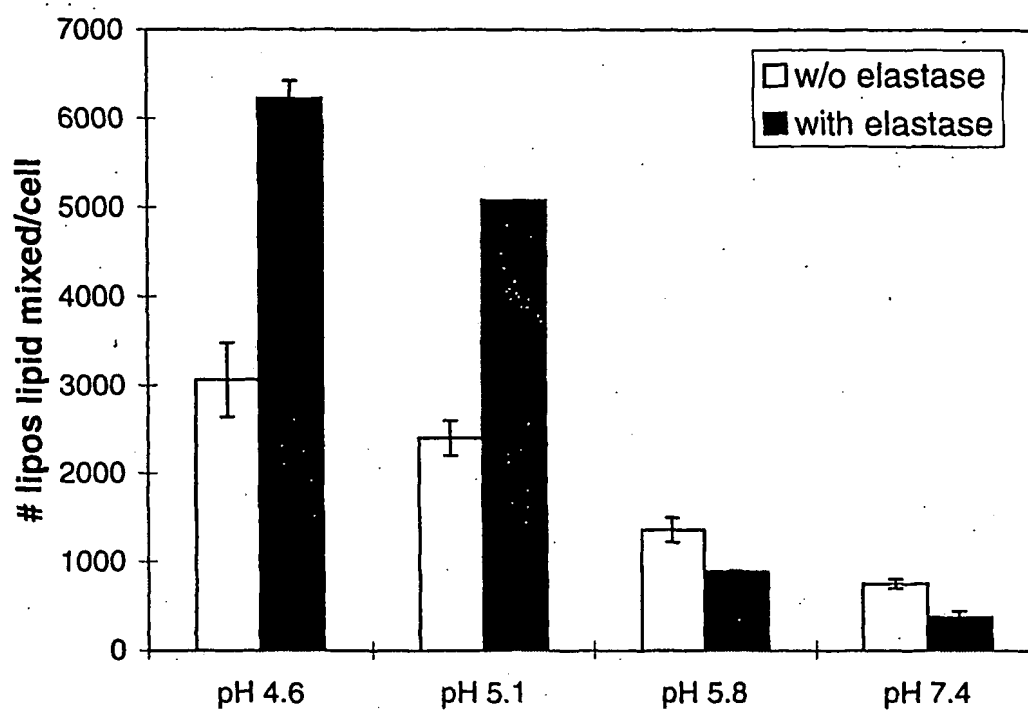
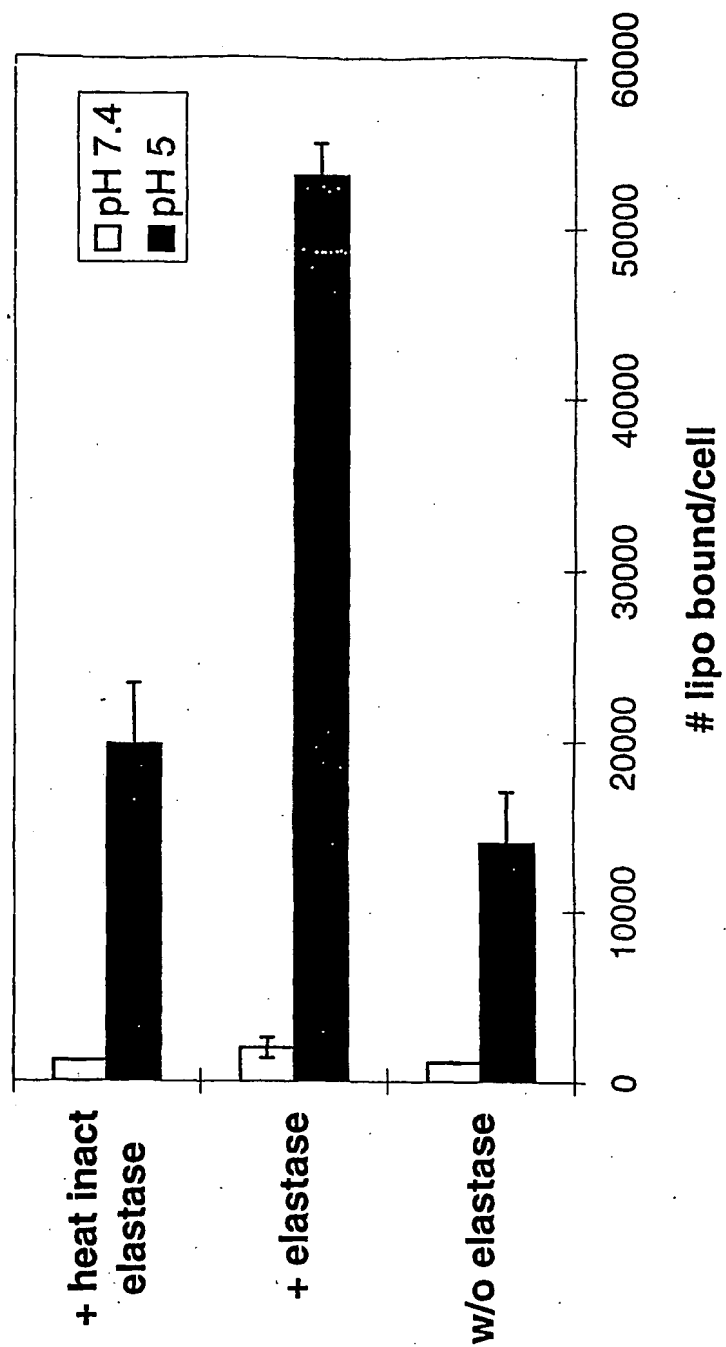


Fig. 16B

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Fig. 17A



2530

Fig. 17B

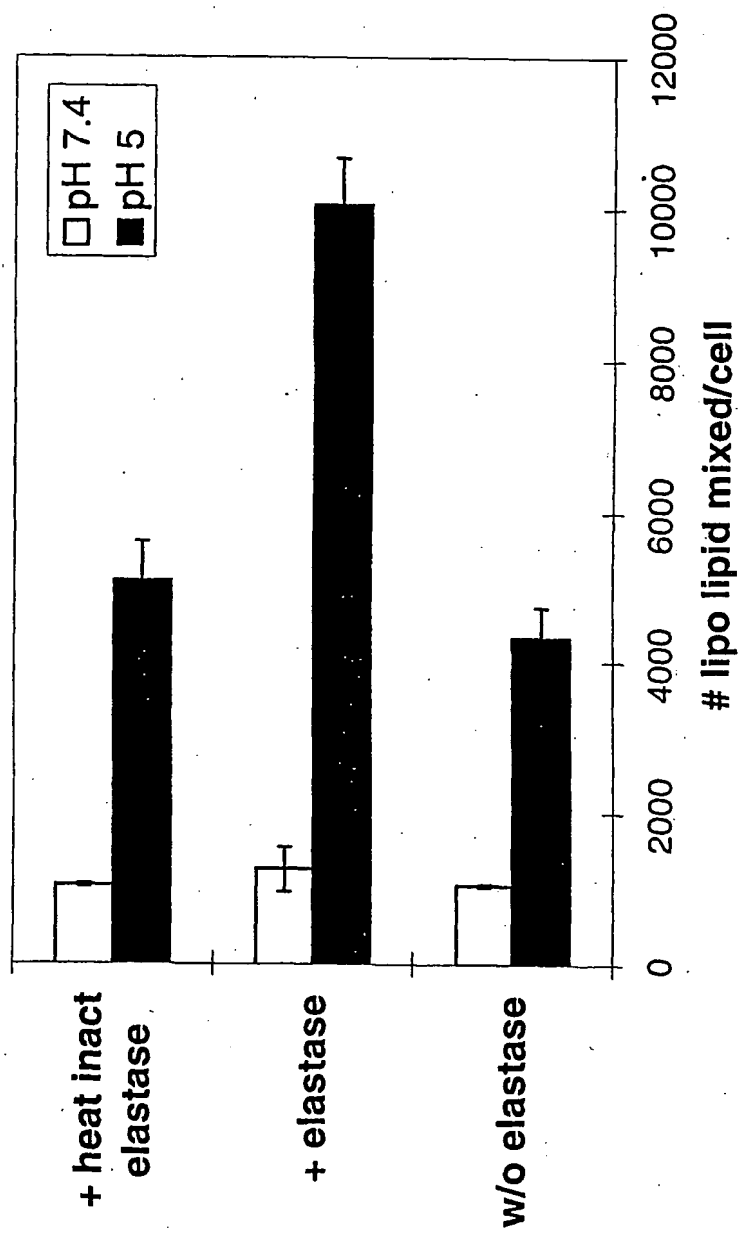


Fig 18A



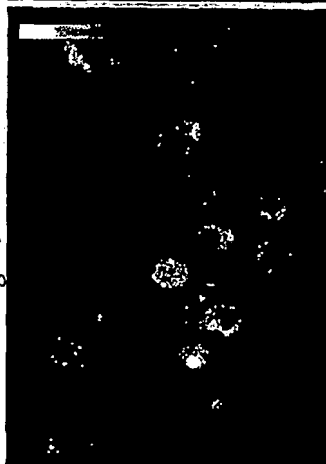
with elastase activation



without elastase activation

Fig 18B

Fig. 19A



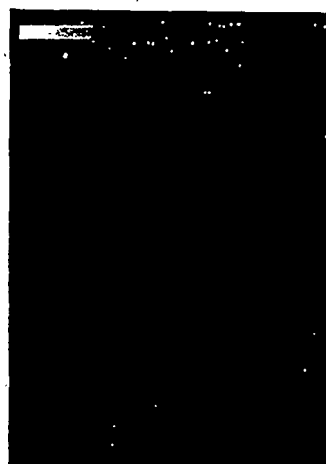
intact lipo, + elastase activation

Fig. 19B



intact lipo, w/o elastase activation

Fig. 19D



freeze/thaw lipo, w/o elastase activation

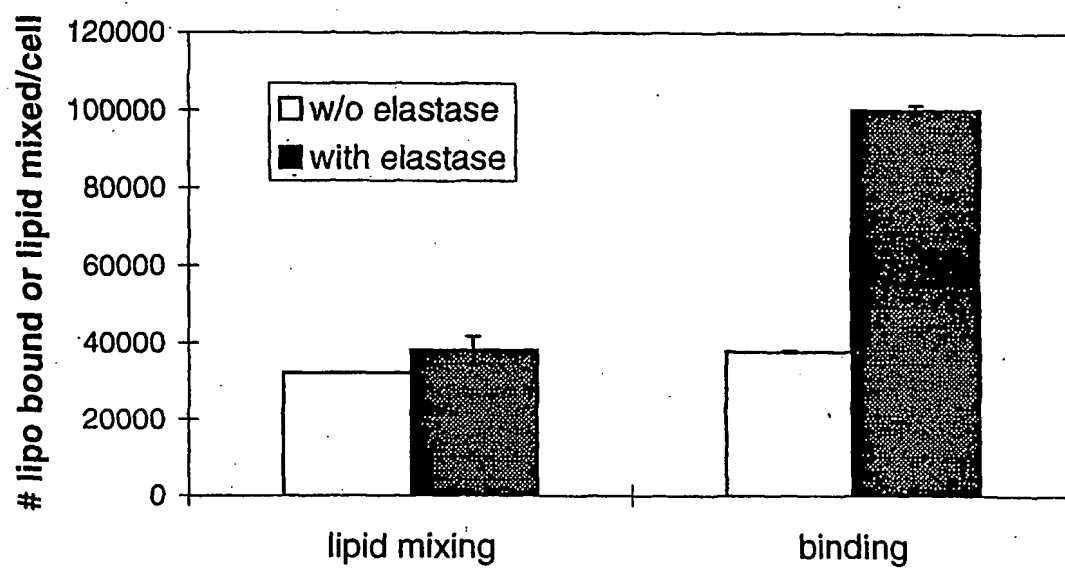
Fig. 19C



freeze/thaw lipo, + elastase activation

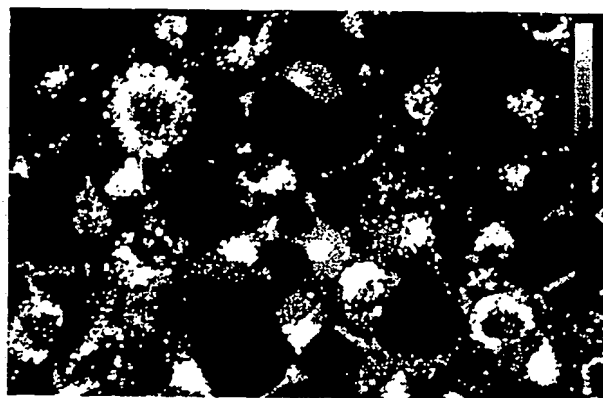
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Fig. 20

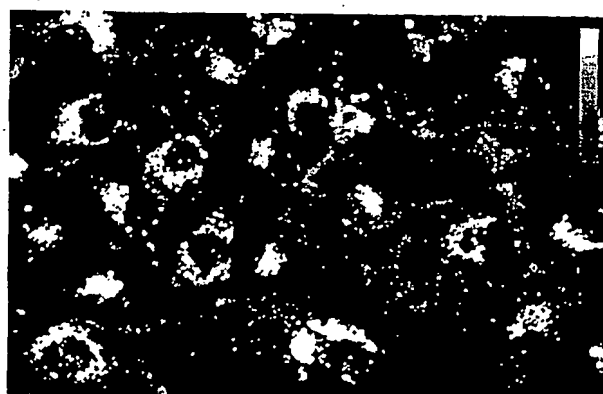


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Fig. 21A



with elastase pretreatment



without elastase pretreatment

Fig. 21B

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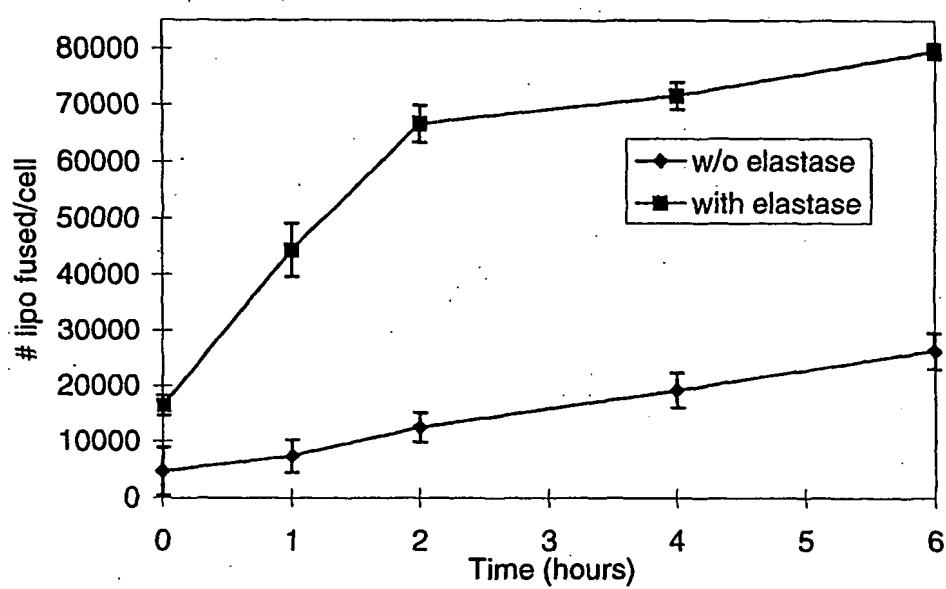


Fig. 22

SEQUENCE LISTING

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 Janoff, Andrew S.
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INTERNATIONAL SEARCH REPORT

International application No.
PCT/US00/16248

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) :A61K 48/00; C12N 15/00, 15/63, 15/88; A61K 9/127

US CL :514/44, 435/320.1, 455, 458; 424/450

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 514/44, 435/320.1, 455, 458; 424/450

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
NONE

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

CAS ONLINE, APS, MEDLINE

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A, E	US 6,087,325 A (MEERS et al) 11 July 2000, entire document, especially columns 39 and 40.	1-26
A	US 5,013,556 A (WOODLE et al) 07 May 1991, entire document.	1-26
A	US 4,837,028 A (ALLEN et al) 06 June 1989, entire document.	1-26



Further documents are listed in the continuation of Box C.



See patent family annex.

* Special categories of cited documents:	*T* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
A document defining the general state of the art which is not considered to be of particular relevance	*X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
E earlier document published on or after the international filing date	*Y* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
L document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*A* document member of the same patent family
O document referring to an oral disclosure, use, exhibition or other means	
P document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search

10 OCTOBER 2000

Date of mailing of the international search report

17 NOV 2000

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